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SUMMARIES


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Patent: WO 02052002-A 132 04-JUL-2002;
Jynavax Technologies Corporation (US)
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Dynavax Technologies Corporation (US)
Location/Qualifiers
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/mol type="genomic DNA"
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NOTE: It currently consists of I contigs. Gaps between the contigs are represented as runs of N. The order of the pieces is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have provided by the submitter. This sequence will be replaced by the finished sequence as soon as it is available and the accession number will be preserved.

* NOTE: This is a "working draft" sequence.

* This sequence will be replaced by the finished sequence as soon as it is this the finished sequence as soon as it is available and the accession number will be preserved.
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Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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synthetic construct
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OJ1136_D07, *** SEQUENCING IN PROGRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (08-AUG-2001) Takuji Sasaki, National Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sasaki, T., Matsumoto, T. and Yamamoto, K. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Published Only in Database (2001)
2 (bases 1 to 119972)
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Patent: WO 02052002-A 14 04-JUL-2002;
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25005 c 25414 g
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Agrobiological Resources, Rice Genome Research Program; Kannondai 2-1-2, Tsukuba, Ibaraki 305-8602, Japan (E-mail:tsasaki@abr.affrc.go.jp, URL:http://rgp.dna.affrc.go.jp/, Tel:81-298-38-7441, Fax:81-298-38-7468)
This PAC clone has 10 kbp sequence gap, which result in 2 contigs (contig a, b). This sequence shows contig b. The orientation of the sequence is from SP6 to T7 of the PAC clone. Genes were predicted from the integrated results of the following:GENSCANI.0, BLASTNI.4, BLASTX2.0 as well as SplicePredictor (October1998 version). The genomic sequence was searched against the non-redundant database NRP(PIR,SWISSPROT, GENPEPT, PDB) from MAFF DNAbank and the cDNA sequence database at RGP. Protein similarities of the cording regions were searched against NRP with BLASTP2.0. ESTs represent the identified cDNA sequences using BLASTNI.4 with the corresponding DDNA accession no. and RGP clone ID.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
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2 (bases 1 to 126038)
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complement(join(1542. 1704 1808. .2005,2133. .2332,2428. .2541,2732. .2868,3006. .3116,5410. .5554))
/note="Similar to glycogenin glucosyltransferase (EC 2.4.1.186). (Z97341)"
                                                                                                                                                                                             CVY1PT1ITLLNSVGTPRSFHLLFFW1LFENVMSLHRTKATL1GLLEAGRANEWVVTE
KLGNALKMKSSSKKSAKKSFMRVWDRLNVTELGVAAFLFSCGWYDLAFGKDHFF1YLF
                                                                                                                                         FQGAAFFIVGIGYVGTIVPQS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /codon_start=1
/protein_id="B
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/mol_type="genomic DNA"
/cultivar="Nipponbare"
/db_xref="taxon:39947"
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note="hypothetical protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         id="BAA82375.1"
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                                                                                         .9351,11606. .11736,12883. .12994,13030. .13164
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ABVINWVMRGVRECVLKADGGCTCSCMKPKLHHLPCSHVLAAAGDCGISPNVYVSNYF
RKEAIFHIWSEEIYGFGISGSYTTLSAQVFYIFDPSKLÆVKKERQTRRIRMDESE
AGMAVDTPALLURGIDBHRSFLSAVBGAQLGTFRFRYRSFEWLÆVDFBHYMLVEBAB
DDRDFAKRWDADRSLLAALVDRWRFPTHTFHLPCGEMAPTLQDVSYLLGLFLAGAAVG
PVDGVFGWKEDLTARFEOVMRLFHLGFTTTLPFYSTVGFSKAWLLQFTTRYAHYQVRR
GYDDFVFETBALQFSTVUMEFYTESAVAARAPCLSSLCTROQAYMLTILPMVFDIF
EPHCFQRVMROFEGLROVFFGNVQFTVLFADHSLITRRGQLAGALWAFRVQQYVDDWYLA
                                                                                                                                                                                   SAARSIMASEVREALESITVYKYSELEKATAGFSEERRVPGTAVYRGVFNGDAAAVKR
VSGDVSGEVGILKKVNHCSLIRLSGLCYHRODYLLVFEYAENGALSWILHGDDAATGV
LGWRQRVQVAEPDVADGLYLHHYNDPCVHKNIKSSNILLDADLHKSSGLARALF
LGWRQRVQVAEPDVADGLYLHHYNDPCVHKNIKSSNILDADLHGVSFGLARALF
AGDGAAAAAQLTRHVVGTQGYLSPEYLEHGLITPKLDVFAFGVVLLELLSGKEAASS
GDGENGEALLLWESAAEALLVDGGGEDAGSNVZAFMDPRLGGDYLJDLAMAVASLAARC
                                                                                                                                                                                                                                                                                                                                            TEEVINELFPHTEENYRDYLRWYLPRTRARVTFTPDAPEPHVAAVTDAYFTHRDRDYF
VARCGRRTSSWSGTTTASRAYRRSRTSTYGTYFARTSSAFERVRRHDRRFHELRRGVR
HLFGRVRQLFLSRSVDPSPTRTGPTSQFYDDDLHGADHHDVLGSSQLGGAPEAHTQEQ
PEVTPVQAGRVGRAVPPDRLTYSHGHIRAQGRRDRGKRPRQ"
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TIAQRNNNTYYDLHTFTSVDDRTKSVLQRAFFSLGACINAFVHCRPVLCIDGTFMTGK
YRGQILTAIGCDGNNQVLPMAFAFVESENTESWYWFLERVHIAVVCMRPNVCLIHDRH
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complement (join(18782. .19019,19135. .19266,19513. .1961

19855. .20113,20190. .20488,21053. .21389,21495. .21645,

22169. .22444,22743. .22774,23219. .24224)

/note="EST C28952(C62945) corresponds to a region of the
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/sranasaasstvvatiavggklpoptacglktvvyaeltagrkavlfa
vpgaftptgsgkhlpoftekapelhakgvda i kcvsvubafvarakksslglgdabvl
llsdgnleltralgvemdlsdkpmglgvrsrryalladdgvvkvlnleeggafttssa
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QCVVQFMETPRVDVEYSLLQQLPLIDTAFLCRKGIDLKKTSRDEHDLRKVTPVYTTIR
CTGHEAVLLLLVLRFGSSLSESLARFARTKVCSDSSECDDDDDDSGDESNS"
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EDFSGLVISEGDHVPWEYKENEVIEGARYAHKDERKEAVKHWAVSLQREFRVVKSTNY
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                                                                                                                               complement (join (32991.
                                                                                                                                                         VARQPAARPAMDEVFVSLAAVYGSTVDWNPSDHGNSGSSLIGR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="5' long terminal repeat"
join(25810. .25825,29004. .31165)
/note="Similar to putative receptor kinase. (AC002332)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complement (25215. .25521)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /protein_id="BAA82378.1"
/db_xref="GI:5441880"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similar to maize transposon MuDR mudrA protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /protein_id="BAA82379.1"
/db_xref="GI:5441881"
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a region of the predicted gene.
Similar to Arabidopsis thaliana BAC genomic sequence.
                                                                                           note="Similar to Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                codon_start=1
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/db_xref="GI:5441879"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similar to Arabidopsis
(AC002292)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /protein_id="BAA82376.1"
/db_xref="GI:5441878"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AC003981) "
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                                   _start=:
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                                                                                                                            .33608,36271. .36273))
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                                                                                              DNA contig
Ś
                                                              Query Match
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Matches 16
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   տ
                                                                 l Similarity
16; Conser
   CGAACGTTCGAGATGA 20
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SdC

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CDS

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/translation="MRRVQLRTHTHARNEIALI PRHTHTRTLGMRLHSYHDTRTHGTG
NVQSTTHAHTCPVPLNVHACI DPGKRRKTPVHSYNMSSSSSPABRPTHATG
TPDRRRSPGSPARAGECRDKGGVVEYRVKEREISPRLVVSSAPRAAAKTSRGGGAAIE
QSRGGGGILGGVTGGNARPKRYATSQIKAEPELLRREVPLTSGKTKKAPKITFKNEPR
HSAPQSDWGTPDSVPBFGPADEYRALRKKYLLLEDENGELDKQLSQAEEEASVLEDEK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /trānslation="maaamattaaaaaeeavvaldketlmlmgcgdappaapcaewet
FKENVRPLKRGRNVGLLNRALKAHADPAQRAALLAARRKMIEAIDEYSGEDPLQPWID
CIKMVQESPPTGGDCSGLVVIYEQCVRAFWHDDRYKNDLRYLKVWLEYAGNCADSEVI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EFPASWSPPKSGGEKLAAAARSRLKRVLSIIQEDA"
complement (join (51409. .51816,53421. .53933))
/note="Similar to Arabidopsis thaliana BAC genomic sequence. (AL021889)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FNEKLNEQTIGVFAVFLSPARGAAPAAAAARSPLKRALSLVEEEA" complement(join(38463. .39039,42006. .42013)) /note="Similar to Arabidopsis thaliana BAC geno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="GI:5441882"
/translation="MLSCIYRPESDIVVESEAPAPPPPPARVIAADGSLRELPASPCV
AVVSSVLGGGDDAASFFUCNSDALYFDDRPPALSPGBPLRPGBLYFVLPRAMLGRPLS
SADMGRLAVRASLALVGERPQRRRRHRQGGGKKQKKVHVNPAQAQSRDDGDGDVDGV
                                                                                                                                                                                                                                                                                                                                                                                                                 join(58631. .58875,59008. .590
/note="Similar to Arabidopsis
sequence. (AC002332)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    predicted gene.
hypothetical protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="Megciasrhagrtesleggepveagsgeasywoygpavislegg
fafcgrgaservaaaaaavflscisrrgspypaaparviaadgslkelhaaaspava
dulrgegssffvcyrsdtlyfnegppamapgealregolyfvlpaamlgoplestadmaa
lavrasaalaavkaperrgarrggdrkmktvrvtplrdegldggdvllheklnertlg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             complement(join(43967. .44533,45832. .4
/note="Similar to Arabidopsis thaliana
fragment. (Z99707)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /trānslation="metcrrrssshqpapaparvitadgslkelavssavadvlrgeg
egrsffvcnsdalyfneqppalapgealrpgqiyfvlpaamlgqplstadmaalavra
saalaaaaakaprkrgvrrgggdrkkktvrvtplrderldggdvllheklnertlgef
                                                                                                                                                                                                                                     FRFLEANQIGQSHTNYYLSYASVMESKNKLKKANEIFNLGIAR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               join(55812. .55816,55861. .55978,56109. .56216,56457. .567:
57319. .57365,57427. .57778)
/note="EST AU069246(C53478) corresponds to a region of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VPAASAADPNSYRRARKGAHRAPERSPFF"
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                                                                                                      /proteIn_id="BAA82386.1"
/db_xref="GI:5441888"
                                                                                                                                                                                                                                                                                                                                        /codon_start=1
/protein_id="BAA82385.1"
/db_xref="GI:5441887"
                                                                                                                                                                                 oin(66733. .66759,67025. .6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proteIn_id="BAA82384.1"
/db_xref="GI:5441886"
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/db_xref="GI:5441884"
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ragment. (299707)
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_xref="GI:5441883"
                        76.2%;
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0;
                        Score 16;
Pred. No.
  Mismatches
                          No.
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25;
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  <u>.</u>
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                                                Length 126038;
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AP005629
LOCUS
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Best Local
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Submitted (07-AUG-2002) Takuji Sasaki, National Institute of Agrobiological Sciences, Rice Genome Research Program; Kannondai 2-1-2, Tsukuba, Ibaraki 305-8602, Japan

(E-mail:tsasaki@nias.affrc.go.jp, URL:http://rgp.dna.affrc.go.jp/, Tel:81-298-38-7441, Fax:81-298-38-7469)

NOTE: It currently consists of 1 contigs. Gaps between the contigs are represented as runs of N. The order of the pieces is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have provided by the submitter. This sequence will be replaced by the finished sequence as soon as it is available and the accession number will be preserved.

* This sequence will be replaced

* This sequence will be replaced

* the accession number will be preserved.

* the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                 33857
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                                         unordered pieces.
AC141727
AC141727.1 GI:29123911
HTG; HTGS PHASE1; HTGS_DRAFT.
Apis mellifera (honeybee)
Apis mellifera
                                                                                                                                                                                                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oryza satīva (japonica cultivar-group) Oryza sativa (japonica cultivar-group) Oryza sativa (japonica cultivar-group) Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa (japonica cultivar-group) chi
OSJNBa0004A19, *** SEQUENCING IN PROGRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sasaki,T., Matsumoto,T. and Katayose,Y. Oryza sativa nipponbare(GA3) genomic DN clone:OSJNBa0004A19
                                                                                                                                                                              AC141727 146568 bp DNA linear HTG Apis mellifera clone CH224-57G3, WORKING DRAFT SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                           5 CGAACGTTCGAGATGA 20
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2 (bases 1 to 144952)
Sasaki,T., Matsumoto,T. and Katayose,Y.
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HTG; HTGS PHASE2.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                 CGAACGTTCGAGATGA 33872
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Oryza sativa
/mol_type="genomic_DNA"
/cultivar="Nipponbare"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="OSJNBa0004A19"
30336 c 30932 g 41
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/chromosome="2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .144952
                                                                                                                                                                                                                                                                                                                                                                                                    76.2%; Score 16; DB
100.0%; Pred. No. 25;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (japonica cultivar-group)"
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ESS ***.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  102 others
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    Apoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RS Mizny, D. M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banke, T., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banke, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Barbaria, J., Bowie, S., Brieva, M., Brown, B., Brown, M., Bryant, N.P., Bulke, T., Chen, G., Chen, G., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox.C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y. Durbin, K.J., Davila, R., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y. Durbin, K.J., Davis, C., Bayer, H., Dugan-Rocha, S., Durbin, K.J., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y. Durbin, K.J., Davis, C., Bayer, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Bhaj, C., Becotto, M., Earnhart, C., Bdgar, D., Edwards, C.C., Bhaj, C., Becotto, M., Earnhart, C., Bdgar, D., Edwards, C.C., Bhaj, C., Becotto, M., Earnhart, C., Bdgar, D., Edwards, C., C., Bhaj, C., Becotto, M., Earnhart, C., Bdgar, D., Edwards, C., Hamilton, K.J., Garza, N., Gill, R., Gabis, A., Gaoli, J., Ford, J., Ford, J., Korat, C., Karlson, E., Halle, S., Hamilton, K.J., Hernandez, J., Homes, J., Jakeson, L.E., Harris, C., Harris, K., Hart, M., Havlask, D., Hawes, A., Hernandez, J., Hernandez, C., Hodgson, A., Hogues, M., Hollows, C., Hodlins, B., Jar, Y., Johnson, R., Jolivet, S., Joudah, S., Karlson, E., Khan, U., King, L., Kovar, C., Kratovic, J., Landry, N., Leevis, L. C., Lewis, L., Lichearge, O., Martin, R., Martindale, A., Martinez, E., Maheshar, Y., Mahabat, K., Morgan, M., Morris, S., Moser, M., Niger, M., Mayen, A., Nigyen, A., Nigyen, A., Nigyen, A., Nigyen, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (19-MAR-2003) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 (bases 1 to 146568)
Worley, K.C.
Direct Submission
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Center: Baylor C
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^{*} NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/cenbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 25 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

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RESULT 9
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/db_xref="taxon:7460"
/clone="CH224-57G3"
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74277: contig of 7530 bp in length
74277: gap of unknown length
81722: contig of 7495 bp in length
81822: gap of unknown length
92875: contig of 11053 bp in length
92975: gap of unknown length
101243: contig of 8268 bp in length
101343: gap of unknown length
113869: contig of 16256 bp in length
113869: gap of unknown length
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12847: gap of unknown length
128547: gap of unknown length
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128547: gap of unknown length
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Oryza satīva (japonica cultivar-group)

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

Enrhartoideae; Oryzeae; Oryza.

El (bases 1 to 166304)

El (bases 1 to 166304)

Chow, T.-Y., Hsing, Y.-I.C., Chen, H.-C., Chen, S.-K., Chen, T.-R.,

Chow, T.-Y., Chang, S.-J., Chen, H.-C., Chen, S.-Y., Hsiao, S.-H.,

Hsiung, J.-N., Hsu, C.-H., Huang, J.-J., Kau, P.-I., Lee, M.-C.,

Leu, H.-L., Li, Y.-F., Lin, S.-J., Lin, Y.-C., Wu, S.-W., Yu, C.-Y.,

Yu, S.-W., Wu, H.-P. and Shaw, J.-F.

Oryza sativa PAC P0681D04 genomic sequence
                                                               Sequence
AR268334
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Chow, T.-Y. and Heing, Y.-I.C.

Direct Submission
Submitted (14-AUG-2002) Institute of Botany, Academia Sinica,
Section 2, Academia Road, Nankang, Taipei 11529, Taiwan
* NOTE: This is a 'working draft' sequence. It currently
* consists of 6 contigs. Gaps between the contigs
* are represented as runs of N. The order of the pieces
* are represented as runs of N. The order of the pieces
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16; Conserva
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166304 bp DNA linear HTG 14-AUG-2002
Oryza sativa (japonica cultivar-group) chromosome 5 clone P0681D04,
*** SEQUENCING IN PROGRESS ***, 6 ordered pieces.
                                        AR268334.1
                                                                                                     AR268334
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                                                                                                                                                                                                           CGAACGTTCGAGATGA 99697
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/mol type="genomic DNA"
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/db xref="taxon:39947"
/chromosome="5"
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8678: contig of 8678 bp in length
8778: gap of unknown length
14903: contig of 6125 bp in length
15003: gap of unknown length
15003: gap of unknown length
1513293: contig of 20390 bp in length
15493: gap of unknown length
15493: gap of unknown length
121124: contig of 85831 bp in length
121124: gap of unknown length
131546: gap of unknown length
13546: gap of unknown length
13546: gap of unknown length
13646: gap of unknown length
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                                                         1 (bases 1 to 22)
Raz, E., Cho, H.J., Richman, D. and Horner, A.A.
Raz, E., Cho, H.J., Richman, D. and Horner, A.A.
Methods for increasing a cytotoxic T lymphocyte
Patent: US 6534062-A 3 18-MAR-2003;
Location/Qualifiers
                                                                                                                                                                  Sequence 3 from patent US AR287743
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Raz, E., Cho,H.J., Richman,D. and Horner,A.A.
Methods for increasing a cytotoxic T lymphocyte response in vivo
Patent: US 6534062-A 1 18-MAR-2003;
Location/Qualifiers
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Patent: US 6498148-A 19 24-DEC-2002;
Location/Qualifiers
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Sequence 1 from patent US
AR308057
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ASSIST PUBL HOPITAUX DE PARIS (FR)
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/db_xref="taxon:32630"
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Dynavax Technologies Corporation (US)
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Methods of modulating an immune response using equences and compositions for use therein patent: WO 011223-A 2 22-FEB-2001;

Dynavax Technologies Corporation (US)
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"

/db_xref="taxon:32630"
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Dynavax Technologies Corporation (US)
Location/Qualifiers
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Technologies Corporation (US)
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/db_xref="ttaxon:32630"
/note="synthetic construct"
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Sequence 2 1
AX250702
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Patent: WO 0168117-A 1 20-SER-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Sequence 1 from Patent WO0168117
AX252291
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Location/Qualifiers
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/db_xref="taxon:32630"
/note="Polynucleotide containing
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/db_xref="taxon:32630"
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AX252509
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Methods of ameliorating symptoms of herpes infection using immunocdulatory polynucleotide sequences

Patent: WO 0168103-A 1 20-SEP-2001;

Pynavax Technologies Corporation (US)

Location/Qualifiers
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Patent: WO 0168117-A 2 20-SEP-2001;
Dynavax Technologies Corporation (U
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/mol type="genomic DNA"
/mol xref="taxon:32630"
/note="Polynucleotide containing
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Methods of ameliorating symptoms of herpes infection using immunocoulatory polynucleotide sequences patent: WO 0168103-A 2 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Dynavax Technologies Corporation (U
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/note="Polynucleotide containing
/note 7 g 6 t
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/mol_type="genomic DNA"
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Immunomodulatory formulations and methods
Patent: WO 0168143-A 1 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="texon:32630"
/note="Polynucleotide containing
4 c 7 g 5 t
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/note="polynucleotide containing
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van Nest,G.
Methods of preventing and treating ng immunomodulatory polynucleotide
                                           synthetic construct synthetic construct artificial sequences.
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Patent: WO 0168116-A 1 20-SBP-2001;
Dynavax Technologies Corporation (US)
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/mol type="genomic DNA"
/mb_xref="taxon:32630"
/note="Polynucleotide containing
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Methods of preventing and treating viral immunodulatory polynucleotide sequences Patent: WO 0168077-A 1 20-SEP-2001;

Dynavax Technologies Corporation (US)
Location/Qualifiers
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Sequence 2 from Patent WO0168077.
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Dynavax Technologies Corporation (US)
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Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="taxon:32630"
/note="Polynucleotide containing
3 c 7 g 6 t
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/mol type="genomic DNA"
/db_xref="taxon:32630"
/note="Polynucleotide containing
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O'Hagan, D., Otten, G., Donnelly, J.J., Polo, J.M., Singh, M., Ulmer, J. and Dubensky, T.W. Microparticles for delivery of the heterologous Patent: WO 0226209-A 19 04-APR-2002; CHIRON CORPORATION (US)
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Immunomodulatory polynucleotides and methods
Patent: WO 02052002-A 2 04-JUL-2002;
Dynavax Technologies Corporation (US)
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/mol type="genomic DNA"
/db_xref="teaxon:32630"
/note="Artificial sequence is synthesized"
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/note="Polynucleotide containing
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Dynavax Technologies Corporation (US)
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Patent: WO 02052002-A 40 04-UUL-2002;
Dynavax Technologies Corporation (US)
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
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Patent: WO 02052002-A 59 04-JUL-2002;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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2 (bases 1 to 22)

2 (Carson,D.A., Raz,E. and Roman,M.

2 (Carson,D.A., Raz,E. and Roman,M.

Immunostimulatory polynucleotide/immunomodulatory mol
Patent: JP 2001503254-A. 34 13-MAR-2001;
THE REGENTS OF THE UNIVERSITY OF CALIFORNIA
OS Artificial Sequence
PN JP 2001503254-A/34
PD 13-MAR-2001
PP 09-CCT-1997 JP 1998518649
PF 09-CCT-1997 UP 1998518649
PF 11-CCT-1996 US 60/028118
PI DENNIS A CARSON, EYAL RAZ, MARK ROMAN
PC A61K39/00,A61K39/385,A61K39/39
CC
FH Key Location/Qualifiers
FT source 1..22
/organism='Artificial Sequence'.
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Patent: WO 03000232-A 1 03-JAN-2003;
Yissum Research Development Company of the Hebrew Univ of Jerusalem (IL); The Regents of the University of California (US)
Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="taxon:32630" 6 t
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/mol type="genomic DNA"
/db_xref="taxon:32630"
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Anti-tumor antigens or their
BD182369
BD182369.1 GI:30793287
WO 02090981-A/1.
                                                                                                                                      Methods of modulating an immune response using immunostimulatory sequences and compositions for use therein
Patent: WO 011223-A 3 22-FEB-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Hanabuchi,S., Ohashi,T. and Kannagi,M.
Anti-tumor antigens or their epitopes against HTLV-1 tumor
Patent: WO 02090981-A 1 14-NOV-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP,SHINO HANABUCHI,TAKASHI OHASHI,
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SHINO HANABUCHI, TAKASHI OHASHI, MARI KANNAGI
G01N13/15, A61X13/10
Description of Artificial Sequence:ISS-ODN
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08-MAY-2001 JP 01P 137526
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/noTe="Synthetic construct"
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/mol_type="genomic DNA"
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Methods of suppressing hepatitis virus infection using immunomodulatory polynucleotide sequences Patent: WO 0168078-A 3 20-SEP-2001;
Dynavax Technologies Corporation (US)
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Dynavax Technologies Corporation (US)
Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="taxon:32630"
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Sequence 3 from Patent W00168144.
AX252522
AX252522.1 GI:15985793
                                                                                                                                                                                                                                        van Nest,G.

Methods of ameliorating symptoms of herpes infection using immunoodulatory polynucleotide sequences
Patent: WO 0168103-A 3 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Patent: WO 0168117-A 3 20-SEP-2001;
Dynavax Technologies Corporation (US)
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/mol_type="genomic DNA"
/mol_tyre="texon:32630"
/db_xref="reaxon:32630"
/note="polynucleotide containing
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Biodegradable immunomodulatory formulations and methods for use
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/db xref="taxon:32630"
/note="Polynucleotide compared to the compared 
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1 (bases 1 to 11033)
Barnett, M.J., Fisher, R.F., Jones, T., Komp, C., Abola, A.P.,
Barloy-Hubler, F., Bowser, L., Capela, D., Galibert, F., Gouz
Gurjal, M., Hong, A., Huizar, L., Hyman, R.W., Kahn, D., Kahn,
Kalman, S., Keating, D.H., Palm, C., Peck, M.C., Surzycki, R.,
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                                                                                                      Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Rhizobiaceae; Sinorhizobium/Ensifer group; Sinorhizobium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Polynucleotide containing
8 c 3 g 6 t
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Pred. No.
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    apela,D., Galibert,F., Gouzy,J.,
Hyman,R.W., Kahn,D., Kahn,M.L.,
C., Peck,M.C., Surzycki,R.,
                                                                                                                                                                                                                                                                    pSymA
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91;
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2 (base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wells,D.H., Yeh,K.-C., Davis,R.W., Federspiel,N.A. and Long,S.R. Nucleotide sequence and predicted functions of the entire Sinorhizobium meliloti psymA megaplasmid proc. Natl. Acad. Sci. U.S.A. 98 (17), 9883-9888 (2001)
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                                                                                                                         /translation="mpadlaartsskifngvdlmdasaradnafyldrogerresnars
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acyfralrddesgginfpaavileavogesgvipapvemlravrfvrbdgiplivde
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vyvdbegldalghfphggsiarmiqhemfragiiletggrfgsvlrlldplvisdae
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                                                                                                                                                                                                                                                                                                                                                                                         /note="glimmer prediction; identical to RhsA (RhbA) from Sinorhizobium meliloti, AADO09412, 0923R2; aminotramsferase class-III-pyridoxal phosphate family; rhizobactin siderophore biosynthesis pathway"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VASPEAPLERRASLEGFERRENAWMILAFALTYRASEGLVRGMEGSYLVDSKVPTEWI
GYMSGAAAATAGLLGALIAALIIRKAGLTATLILLGGLRSLCELAFALNAFGIWPGIA
VAMSASAFQTLIRYMELVAIYSFFMASSSDDQPGTDFTILSCAELVVYLIGTSIAGYV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /translation="MLAAVVQGSDPIMTIAQTSPAVREGSTAAGAGRLYAVLGGLYLA
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TLAIGNAVQAGSIALGVIVGGTLTLVLFHKIGWRPTILLVACLSLLPLVAAIWMKDRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Irp8 from Yersinia enterocolitica,
/codon start=1
/transI_table=11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product="putative transmembrane transport protein"
/protein_id="AAK65915.1"
/db_xref="GI:14524428"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /function="Cell processes; Transport of small molecules"
/note="glimmer prediction; similar to signal transducer
Trp8 from Yersinia enterocolitica, CAB46571"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /gene="SMa2337"
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                                                                                                                                                                                                                                                                                                                          product="RhbA diaminobutyrate-pyruvate aminotransferase"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |522. .2934
|gene="rhbA"
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                                                                                                                                                                                                                                                                                    protein_id="AAK65916.1"
db_xref="GI:14524429"
                                                                                                                                                                                                                                                                                                                                                     codon_start=1/
transl_table=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ncorporation metal ions"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene="rhbA"
EC_number="2.6.1.46"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        strain="1021"
                                                                     gene="rhbB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            function="Cell processes; Transport of small molecules;
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                                              note="synonym:
                                                                                                          dqvsgalaaaferlgrkaa"
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, USA
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DILNR?HSSRLI BAALHARNODLAALAGDEVSFVAAROGLI ACHG HIPCPKSREGMT
EAESRRYSPEFAAGFPLRWFAVESELFHTGHSQOSPSAEWLKRAMGSDI DALKARPI
EAESRRYSPEFAAGFPLRWFAVESELFHTGHSQOSPSAEWLKRAMGSDI DALKAPL
AGDESILLPVHFWQADOMLKDFTVAALVAAGRWLDCGEAGKFWFPTSSVSTLYREDASF
MLKMSLGVGI TNS VRVNLARELLRGDDMYRFRRHELWQDFSRSYPGLTLI PDPAFMGV
KIDGALI DGLS VSMRENFTGANADRNVSLLAAVCHL PDRGSELGALI RNRAHLERR
PLDI VARDWFERFFLTL FYRF F FGLYLLHGI DAWBAHOONI VVELEHGYP TGLYFXDNOG
FFHHERAHGALVEALPGFGEPSES VFGEEPVDERLLYYAFINSVLGMVGALGREGLVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VDQTAAASSLYSGTSFQGLRGLIDDLDPLPEVGTGIAAALAEIGRPALEHAMVVGHPA
AMAHLHCPVAVPALAAEVLISATNQSLDSWDQSPPATLVEERVLACLTQLAELPASAS
GNFTSGGTQSNMTALYLAAVRCGPDARKAGVVLTSAHAHFSIRKSAAILGFAEDAVIA
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PAEKKDRGWHMLVGEBSFFGRGIAPANIRAFTRFLFLDDPGTQKVVGEBSVAARRLLR
                                                                                                                                                                                                                                                                                                                                                                                                                                                          /function="Cell processes; Transport of small molecules; incorporation metal ions" /note="glimmer prediction; identical to RhsD (RhbD) from Sinorhizobium meliloti, Q923Q9, AFI10737"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WMHVDAAYGGGLLFSRHRSRLEGLEHAHSITLDFHKMLFQPISCGVLLLRDRADFAPL
ASKADYLNPEDAVFADAPNLVERSMQTTRRADALKILMTMRAIGRDGLDALICQTLQN
THAAAEAVKTREYLSLAGPPSLSTVLFRYVSARGPKFADAITLKTRAALFNAGIAALA
                                                           /note="synonym: 6742. .8112
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GGSLGLPAHYRSATGHHLFGEFVMLTDENGAKTVDAVEAISAVIERLEFSIAAKDGRV
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                                                                                                                                                                                                                                                                                                                                                     protein"
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/db_xref="GI:14524431"
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/transI_table=11
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/note="synonym:
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/function="Cell
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                                                                                                                                                                                                                                                                                                                                                                                                           trans1
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function="Cell proce
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processes; Transport of small molecules;
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ORGANISM
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AF110737
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                                                              REFERENCE
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2 (bases 1 to 11853)
Lynch,D., O Connell,M. and
Cloning and sequence analys
                                                                                                                                                Genetic organization of the region encoding regulation, biosynthesis, and transport of rhizobactin 1021, a side produced by Sinorhizobium mellioti
J. Bacteriol. 183 (8), 2576-2585 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                 Sinorhizobium meliloti (Rhizobium meliloti)
Sinorhizobium meliloti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AF110737 11853 bp DNA linear BCT Sinorhizobium meliloti strain 2011 rhizobactin regulon,
                                                                                                                                                                                                                                                                       and O'Connell, M.
                                                                                                                                                                                                                                                                                                                                                       Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales; Rhizobiaceae; Sinorhizobium/Ensifer group; Sinorhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF110737.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AF110737
                                                                                                                                                                                                                                                                                          ynch,D., O'Brien,J., Welch,T., Clarke,P., Cuiv,P.O., Crosa,J.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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CAKLAARGGSVDTHYMMPVHPMOWDHMIVPHFAADIAAGHIVFLGKGDDLYLPQOSVR
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ETFHHGVGAPDLGLGAFRNAVIVNQLLGREHYRVNASASFQKFGLPSSQTAPSSISGD
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/note="glimmer prediction; identical to RhsE (RhbE) from
Sinorhizobium meliloti, Q9Z3Q8, AAD09416"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TLSNI SHPEKSTLKLCMTI LNTAVYRG I PGKRALTAAPLTTWLDQLLARDQFLSEECG
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Sinorhizobium meliloti, 09Z307, AAD09417"
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/transI_table=11
/product="RhbF Rhizobactin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="synonym: SMa2410"
3096. .9901
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transl_table=11
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ncorporation metal ions"
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   M. and O Brien, J.
analysis of the Sinorhizobium meliloti 2011
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Submitted (03-DEC-1998) School of Biotechnology, Dublin City University, Glasnevin, Dublin, Leinster Dublin 9, Ireland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University, Glasnevin, Dublin Location/Qualifiers
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Lynch,D., O Connell,M. and O Brien,J.
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42. .1454
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/strain="2011"
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/codon start=1
/transI_table=11
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/function="siderophore biosynthesis"
/note="similar to DABA decarboxylase"
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note="FUR Binding Site"
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                                                                                                                                                                                                                                                                                                  /gene="rhbC"
/function="siderophore biosynthesis"
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DILINRTHSSRLLIEAALHARNGDLAALAGDEVSFVAAEQGLIAGHGIHPCPKSREGMT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      function="siderophore biosynthesis" note="similar to IucB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note="previously rhsD'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene="rhbD"
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COMPLEMENT (8495. .9439)
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complement (8495.

.9439)

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JOURNAL
MEDLINE
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REFERENCE
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AF381613
LOCUS
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TITLE
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                                                                                                2 (bases 1 to 18662)
Jones, B.F. and Hotez, P.J.
Direct Submission
Submitted (17-MAY-2001) Department of Epidemiology and Public Health, Yale University School of Medicine, 60 College St., Ne Haven, CT 06510, USA
                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida; Ancylostomatoidea; Ancylostomatidae; Ancylostomatinae; Ancylostoma.

1 (bases 1 to 18662)

Jones, B.F. and Hotez, P.J.

Molecular cloning and characterization of Ac-mep-1, a developmentally regulated gut luminal metalloendopeptidase from adult Ancylostoma caninum hookworms

Mol. Biochem. Parasitol. 119 (1), 107-116 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AF381613.1
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RADVIFLPSDASSEITLPEGGRFDCAHLPAYALASKRDLLKPIMMQPLAAECLFLQLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGGRSAQQISEIARTIYVVDSDQIQAEARSGKTLQQILGETIPSFDPASDGARTSFGQ
NLRGRPPLILVDGVSMNSARSLSRQFDAIDPFNIERVEVLSGATAIYGGNATGGIINI
ITKKGKDAEPGLHAEVTGGMGSGFAGSQDFDRNAAGAVTYNSENWDARLSIAGNRTGA
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                                                           Location/Qualifiers
j. .18662
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NLANGSFSITDVAEAERITPRAIQKFFSREGTTFSRYVLGRRLSLAKSLILAEGEATS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene="rhizobactin regulon"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "SQIAYNVGFNDLSYFNRTFRSRYGVRPSDLRRLAAAA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GI:14549936
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Direct Submission
Submitted (16-NOV-1999) Celera Genomics, 45 West Gude Drive,
Rockville, MD, USA
Phis sequence was identified as CDM:10210347 by the submitter.
For further information on this sequence e-mail to fly@celera.com.
* NOTE: This is a 'working draft' sequence.
* This sequence will be replaced
* the finished sequence as soon as it is available and
* the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                              AC013960.1 GI:6437375
HTG; HTGS PHASE2.
Drosophila melanogaster (fruit fly)
Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 83321)
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join(12956. .12983,13860. .14239,15228. .15300,15357. .1
15607. .15689,15756. .15914,16878. .17031,17116. .17184,
17252. .17380,17443. .17544,17599. .17696,17963. .18060,
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15607. .15689,15756. .15914,16878. .17031,17116. .17184,
17252. .17380,17443. .17544,17599. .17696,17963. .18080,
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                                                                                                                                      Location/Qualifiers
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protein_id="AAK67057.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene="MEP-1"
codon_start=1
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100.0%; Pred. No. 1e
Live 0; Mismatches
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r. *** SEQUENCING
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Swith, H.O., Rubhn, G.M. and Venter, J.C.,
Sequencing of Drosophila chromosome 3R, region 83A-83B
Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence was assembled using end sequences from a whole genome shotgun and from subclones of this BAC and its neighboring clones. For further information about this sequence, including its location and relationship to other sequences, please visit our sequence archive web site (http://www.fruitfly.org/sequence/) or send email to bdgp@fruitfly.berkeley.edu.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (17-AUG-1999) Drosophila Genome Center, Lawrence Laboratory, MS 64-121, Berkeley, CA 94720, USA On Feb 22, 2001 this sequence version replaced gi:7143384.
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Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 168448)
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Berkeley Drosophila Genome Project
Lawrence Berkeley National Laboratory,
Berkeley, CA 94720
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
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3B, BAC clone
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TITLE
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bentley, S.D., Chater, K.F., Cerdeno-Tarraga, A.M., Challis, G.L., Thomson, N.R., James, K.D., Harris, D.E., Quall, M.A., Kieser, H., Harper, D., Bateman, A., Brown, S., Chandra, G., Chen, C.W., Collins, M., Cronin, A., Fraser, A., Goble, A., Hidalgo, J., Hornsby, T., Howarth, S., Ruben, C.H., Kieser, T., Larke, L., Murphy, L., Oliver, K., O'Neil, S., Rabbinowitsch, E., Rajandream, M.A., Rutherford, K., Rutter, S., Seeger, K., Saunders, D., Sharp, S., Squares, R., Squares, S., Squares, S., Squares, S., Squares, S., Squares, S., Parkhill, J. and Hopwood, D.A., Parkhill, J. and Hopwood, D.A., Complete genome sequence of the model actinomycete Streptomyces coelicolor 33(2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             295150 bp DNA linear BCT 11-FEB-2003
Streptomyces coelicolor A3(2) complete genome; segment 23/29.
AL939126 AL009199 AL023517 AL023702 AL031514 AL034446 AL034492
AL035212 AL035478 AL049727 AL132824 AL132997 AL589708 AL591522
AL645882
AL939126.1 GI:24413894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (09-MAY-2002) Submitted on behalf of the Streptomyces sequencing team, Sanger Institute, Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 15A E-mail: sdb@sanger.ac.uk On or before Oct 26, 2002 this sequence version replaced gi:20520744, gi:20520662, gi:20520663, gi:20520667, gi:20520669, gi:20520761, gi:20520816, gi:2052067, gi:20520820, gi:20520846, gi:20520823, gi:20520883, gi:20520869.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Streptomyces coelicolor A3(2)
Streptomyces coelicolor A3(2)
Bacteria, Actinobacteria, Actinobacteridae; Actinomycetales;
Streptomycineae; Streptomycetaceae; Streptomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bentley, S.D.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 295150)
                                                                                                                                                                                                                            /note="SC7B7.02, possible lipoprotein, len: 469; has similarity to signal sequence of Mycobacterium legrae hypothetical protein TR.E334742 (EMBL.238604) MLCB2052.27 (445 aa), fasta scores; opt: 75 z-score: 199.2 E(): 0.00062, 23.3% identity in 335 aa overlap. Contains N-terminal signal sequence and appropriately positioned PS00013 Prokaryotic membrane lipoprotein lipid attachment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /gene="SC06005"
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                                                                                                                                                                                                                                                                                                                                                                                                               gene="SC06005"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               note="synonym:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            db_xref="taxon:100226"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .295150
                                                                                                                                                                                     start=1
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                                                                                                     misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2589.
                             inner membrane
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NTRPGLILTYVAFALFFTMFFLYSFFRSLPHDVYEAALIDGAGDWRAFFQVMLPMARP
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                                                                                                                                                                                                                                                                                                                                                                                  /note="SC707.04, probable integral membrane transport /note="SC707.04, probable integral membrane transport protein, len: 301 aa; similar to many members of the MAL subfamily of the binding-protein-dependant transport systems membrane proteins eg. TR:Q06111 (EMBL:X6092) putative transport system permease protein from Clostridium perfringens (275 aa), fasta scores; opt: 590 Z-score: 796.5 E(): 0, 33.6% identity in 265 aa overlap. Contains E300402 Binding-protein-dependent transport systems inner membrane comp signature"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LAIFVI FVLI PEGQAI FWGMTDWRGFS PDYNFVGFDNFTKMFQDDI FLKALRNVALLA AFVPLVTLTILALGVAVAI TILGGPSKGPVRGI IGAS FVRI I SFFPYVVPAI I VUGLIWAQ MYDPNAGLLNGVLTGLGDPSKGPVRGI IGAS FVRI I SFFPYVVPAI I VUGLIWAQ MYDPNAGLLNGVLTGLGDPDTFANLGEKAAAMPAVMFVI WGLVGFYAVLFI AAI K GVPGELYEAAKI DGAGRFRTTI SITLIPAI RDSVQTAVI YLGI AALDAFVYQQAMVPNG GPDNSTLTI SORLFNVAFAKQOFGYATAMGVVLAAVTLVFAALVFLVNRLTGGGEGES KKKAPGSRARRAAAKGGAR"
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fasta scores; opt: 393 z-score: 636.8 E(): 2.6e-28, 32.5%
identity in 280 aa overlap. Contains PS00402
Binding-protein-dependent transport systems inner membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein, len: 353 aa; similar to many members of the MALFG subfamily of the binding-protein-dependant transport system membrane proteins eg. MSMF_STRMU Q00750 multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HGKEAATYYRTLLIDSAIKEGGDEVRLIALENLEKGCWSHPAVQGVIKVMETMVKQKMF
VPGGSGOTQFQKAQAIWSNDQKALLYPSGGWIENEMKKATKADFQMTGIESMTLTDKPA
LPYEALRAAAGEPFIYMKQGKWPAGGKEVLRAMLSEKAAANESKTKLAPTIVKGTVPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="synonym: SC7b7.04"
2999. .3907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'gene="SC06006"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /product="probable trans
/protein_id="CAA15783.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'gene="SC06007"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene="SC06006"
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transl_table=11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene="SC06006"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note="PS00013 Prokaryotic membrane lipoprotein lipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note="possible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     translation="MKDT1PTAETASRRPEPAARGGRPRRRKLTFDRVTFFLAFLGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Binding-protein-dependent component signature"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transport systems
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Binding-protein-dependent transport systems comp signature"

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RESULT
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Best Local
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62
                                                                      17025
                                                                                                                                                                         15;
                                                                                                                                                                                                Similarity
                                                                      TCGTCGAACGTTCGA 17039
                                                                                                                     TCGTCGAACGTTCGA 15
                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                            /translation="mntrmrraavavattamavslaacgsakesdggsdssdsaakkg ddikycillebnotary ekepdkemiekkykei.tnnkgevyavakkgdaslonoovdtm vtnkvdymiidavdykalagsvkkahdagikkei.tnnkgevyavakgdaslonoovdykei.tnnkoevyavakgdbeldaytsednyvgktykkendagkevdykentidaytsednytykekye geallikakgdgkkigkesddtkekkpe geallikakgdgkei.talkragiadipytgdbaelagvor ivtgegymsveksypkeaetaaemavalakgesitalkragiadipytgdpaelagvor ivtgegymsveksypkeaetaaemavalakgesidsindkydsattkgvpavivpvv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein precursor (330 aa), fasta scores; opt: 619 z-score: 573.4 E(): 8.9e-25, 35.7% identity in 325 aa overlap. Contains N-terminal signal sequence and appropriately positioned PS00013 Prokaryotic membrane lipoprotein lipid attachment site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="SC7B7.06, probable solute-binding protein of transmembrane transport system, len: 403 aa; similar to many eg. XYLF_ECOLI_P37387 d-xylose-binding periplasmic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5475
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LAEESEFLDVDASSDQGFDRAEQLVSRLIEATGVDRAKIAFVGLGVFGPIDVESGTLG
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HGTDLTWEGVVRLARDGDPGCRRVIADVGRHIGSGVANLCNLLNPSRVVLGGDLAEAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="SC7B7.05, probable transcriptional repressor protein, len 403 aa; similar to many eg. XYLR ANATH Q44406 xylose repressor from Anaerocellum thermophilm (39 aa), fasta scores; opt: 548 z-score: 767.3 E(): 0, 33.1% identity in 393 aa overlap. Contains PS01125 ROK family signature and helix-turn-helix motif at aa positions 29-50
                                                                                                                                                                                                                                                                                                                                                     SLTKDNIKETVIKDGFYTIDEICTDKYKAACDKIGLK" 5514. .5546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="hairpin loop with 23bp stem"
4128. .4131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GALPAATPAFT"
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4142. .5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /gene="SCO6008"
/note="PS01125 ROK family signature"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /product="probable transcriptional repressor protein"
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/note="PS00013 Prokaryotic membrane
attachment site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /product="solute-binding protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'gene="SC06009"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /codon_start=1
/transl_table=
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                                                                                                                                                                    71.4%; Score 15; DB 1; Lo 100.0%; Pred. No. 1.1e+02; ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Addms, M.D., Celniker, S.E., Holt, R.A., Evans, C.A., Gocayne, J.D., Andams, M.D., Celniker, S.E., Li, P.W., Hookins, R.A., Galle, R.F., George, R.A., Lewis, S.E., Richards, S.A., Ashburner, M., Henderson, S.N., Sutton, G.G., Wortman, J.R., Yandell, M.D., Zhang, Q., Chen, L.X., Button, G.G., Wortman, J.R., Yandell, M.D., Zhang, Q., Chen, L.X., Brandon, R.C., Rogers, M.H., Blazej, R.G., Champe, M., Pfeiffer, B.D., Wan, K.H., Doyle, C., Baxter, B.G., Helt, G., Natzer, B.G., Champe, M., Pfeiffer, B.D., Wan, K.H., Doyle, C., Baxter, B.G., Helt, G., Champe, M., Pfeiffer, B.D., Barllew, R.M., Basu, A., Baxendale, J., Bayraktaroglu, L., Beasley, E.M., Beeson, K.Y., Benos, P.V., Berman, B.P., Bhandari, D., Botshakov, S., Borkova, D., Botchan, M.R., Bouck, J., Brokstein, P., Brottier, P., Bartley, R.C., Busan, D.A., Butler, H., Cadleu, E., Center, A., D., Dew, J., D., Botchan, M.R., Bouck, J., Brokstein, P., Brottier, P., Bartley, R.C., Davies, P., John, R., Butler, H., Cadleu, E., Center, A., D., Dew, J., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., Dietz, S.M., Dodson, K., Doup, L.E., Downes, M., Dugan-Rocha, S., College, A., Gong, F., Gorrell, J.H., Gu, Z., Gelbart, R., M., L., Harvey, D., Helman, T.J., Gu, Z., Gelbart, R.M., Harris, M.L., Harvey, D., Helman, T.J., Gu, Z., Gelbart, R.M., Martis, M.L., Harvey, D., Helman, T.J., Hernandez, J.R., Mouk, J., Hoetin, D., Hostin, D., Karft, C., Kravitz, S., Kulp, D., L., Nelson, D., Karshilov, G., Milshina, N.V., Mobarry, C., Morry, D., Martis, M., Kalush, F., Karshi, C., Kravitz, S., Kulp, D., Scheder, F., Shen, H., Shue, B.C., Siden-Kiamos, I., Stapleton, M., Sto
                                                                        Celniker, S.E., Adams, M.D., Kronmiller, B., Wan, K.H., Holt, R.A., Evans, C.A., Gocayne, J.D., Amanatides, P.G., Brandon, R.C., Rogers, Y., Barzon, J., An, H., Baldwin, D., Banzon, J., Beeson, K.Y., Busam, D.A., Carlson, J.W., Center, A., Champe, M., Davenport, L.B., Dietz, S.M., Dodson, K., Dorsett, V., Doup, L.E., Doyle, C., Dresnek, D., Farfan, D., Ferriera, S., Frise, E., Galle, R.F., Garg, M.S., George, R.A., Gonzalez, M., Houck, J., Hoskins, R.A., Hostin, D., Howland, T.J., Lbegwam, C., Jalali, M., Kruse, D., Li, P., Mattei, B., Moshrefi, A., McIntosh, T.C., Moy, M., Murphy, B., Nelson, C., Nelson, K.A., Nunoo, J., Pacleb, J., Paragas, V., Park, S., Patel, S., Pfeiffer, B., Phouanenavong, S., Pittman, G.S., Puri, V., Richards, S., Scheeler, F., Stapleton, M., Strong, R., Svirekas, R., Tector, C., Tyler, D., Williams, S.M., Zaveri, J.S., Smith, H.O., Venter, J.C. and Rubin, G.M.
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3 (bases 1 to 296091)
Misra, S., Crosby, M.A., Mat
Campbell, K., Hradecky, P.,

Matthews,B.B., Bayraktaroglu,L., P., Huang,Y., Kaminker,J.S., Prochnik,S.E.,

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AUTHORS
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Location/qualifiers
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Annotation of Drosophila melanogaster genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (06-SEP-2002) University of California Berkeley, 539 Life Sciences Addition, Berkeley, CA 94720, USA 6 (bases 1 to 296091)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (21-MAR-2000) Celera Genomics, Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
4 (bases 1 to 296091)
4 (bases 1 to 296091)
Adams, M.D., Celniker, S.E., Gibbs, R.A., Rubin, G.M. and Venter, C.J.
Direct Submission
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complement/~
                                                                                                                      QYCRSMRDQILDSFSHRDFSIIVVGNKFDNVTBAQANSQQTFTQYRILSITAHPCHHT
YPELAEWQELKDISTLVRKHWRCGYVECSAQYNYKIGDVFRELMGCTSAVGGGGAGGG
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13991. .15004
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chromosome="3R"
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%; Pred. No. 1.1
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15; Conserv
Immunomodulatory polynucleotides and methods of using the same Patent: WO 02052002-A 98 04-JUL-2002; Dynavax Technologies Corporation (US)
                                                                                                                                                                  synthetic construct
synthetic construct
                                                                                                                                                                                                                                                                       Sequence 98
AX592408
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                                                                                   Fearon, K.L. and Dina, D.
                                                                                                                                      rtificial sequences.
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RLDGVI WTGAEERAASICH WATTFAE JANARENTVSSVFSCLAVHACHLDGVI
RHKLFTKCFCVFAQTKTRHPELMODURFTFRI PHARMNEVQEEDLSDCGYSVLTWSAE
AGVDCFVKQQKNSLFVHFQGHEEYETQSLLGEYRDMGRFLRGENEVCFTI PRGYLNK
GAEEILI AFRQKALSDRQPELFADFPADQLTRDLSNVMHLPAQHIYRNWLQYMVSQPA
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/gene="bl16280"
/note="ORF_ID:bl16280
unknown protein"
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/gene="bl16280"
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Dynavax Technologies Corporation (US)
Location/Qualifiers
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Sequence 29 from Patent WO02052002.
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                   /note="Polynucleotide containing
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Sequence 44 from Patent WO02052002.
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Patent: WO 02052002-A 44 04-JUL-2002;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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                                                                 /organism="synthetic construct"
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/noFe="Polynucleotide containing
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Patent: WO 02052002-A 55 04-JUL-2002;
Dynavax Technologies Corporation (US)
Location/Qualifiers
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Sequence 55 from Patent W002052002.
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/mol type="genomic DNA"
/mol xref="taxon:32630"
/note="Polynucleotide containing
4 c 7 g 5 t
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Use of cationic lipids for intracellular protein delivery
Patent: WO 0143778-A 1 21-UUN-2001;
Gene Therapy Systems, Inc. (US)
Location/Qualifiers
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Sequence 1 from Patent W00143778
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Sequence 9 from Patent WO0168103
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                                                               /organism="synthetic construct"
/mol type="genomic DNA"
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/note="Polynucleotide containing
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f c 6 g 8 t
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/db_xref="texon:32630"
/note="Polynucleotide containin
6 c 6 g 8 t
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LOCUS
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Search completed: December 17, 2003, 09:38:42 Job time: 1236 secs
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Matches 14; Conserv
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                                                                                         248 ACGTTCGAGATGAT 261
                                                                                                                                                                         14; Conservative
                                                                                                             8 ACGTTCGAGATGAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (01-SEP-1999) Genoscope - Centre National de Sequencage: CP 5706 91057 EVRY cedex - FRANCE (B-mail: seqref@genoscope.cns.fr - Web: www.genoscope.cns.fr)
The cDNA library to be analyzed within the framework of this project was created using a Botrytis cinerea strain which was grown under conditions of nitrogen deprivation, which is the normal situation for B. cinerea during its development on its host plant. The library was produced in an oriented direction, in the pBSII
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CDNA library; nitrogen deprivation.

cDNA library; nitrogen deprivation.

Botryotinia fuckeliana

Botryotinia fuckeliana

Eukaryota; Fungi; Ascomycota; Pezizomycotina; Leotiomycetes;

Helotiales; Sclerotiniaceae; Botryotinia.

1 (bases 1 to 600)
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Botrytis cinerea strain T4 cDNA library under conditions of
nitrogen deprivation.
AL110950
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Direct Submission
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                                                                                                                                                                                                                                                                                    189
                                                                                                                                                                                                                                                                                                                                            /organism="Botryotinia fuckeliana"
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103 c 137 g 171 t
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                                                                                                                                                                  66.7%; Score 14; DB 8; Length 600
100.0%; Pred. No. 4.1e+02;
tive 0; Mismatches 0; Indels
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Perfect score:
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                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                              Score
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seq length: 2000000000
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Gapop 60.0 , Gapext 60.0
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AA097939
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SX455352
CC235774
CNS09N54
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CNS09Q90
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BM624679
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BM645614
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TA99D10P
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BM632552
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BM866297
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AQ623639 HS_5377_A
BHB16458 AW_Ba007
BZ391656 BINCQ64TR
CD375545 PTMM00709
BX455352 BX455352
CC235774 CH261-12
AA186214 T3860 MA
BX070096 Single re
R04873 pk.33h10 NL
AA186214 T3860 MA
BX070096 Single re
B1597844 BB170008B
B1590565 BB170006874
AQ252378 HS_2262
AA186214 T3860 MB
B1590565 BB170006874
AQ252378 HS_2262
AA20752378 HS_2262
AA302528 Cta313b12.
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BM659639355 170006874
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BM6597586 AL697586
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JOURNAL COMMENT
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BU095678/c
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              source
                                                                                                                                                                                                                                                                                                                                                                                         sequence.
BU095678
                                                                                                                                        Email: dsanchez@ilb.unsam.edu.ar
Email: dsanchez@ilb.unsam.edu.ar
Sequences were basecalled with phred and vector was masked with
crossmatch (see http://www.phrap.org). Sequences were then trimmed
from both ends to remove low quality bases and masked vector.
                                                                                                                                                                                                                  Genomics and Bioinformatics
Instituto de Investigaciones Biotecnologicas
Av. Gral Paz S/N, INTI, Edificio 24, B 1650 KNA,
                                                                                                                                                                                                                                                                      genes
                                                                                                                                                                                                                                                                                carassii: identification of trans-sialidase-like
                                                                                                                                                                                                                                                                                        Aguero, F., Campo, V., Cremona, L., Jager, A., Sanchez, D.O. and Frasch, A.C. Gene discovery in the freshwater fish paras
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EST.
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tca-163 tca Trypanosoma
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Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                      from both ends to
Plate: 01 row: n
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                                                                                                                                                                                                                                                                                                                              rypanosoma.
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                                                                                                                             primer: T7
                                                                                                                                                                                      (54-11) 4580/7255/7
(54-11) 4752-9639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Argentina
/dev_stage="blood trypomastigote"
/lab_host="Goldfish (Carassius auratus)"
/clone_lib="cca"
/note="Vector: pSport1; Blood trypomastigotes were
/note="Vector: pSport1; Blood trypomastigotes were
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et al. Parasitol Res (1998) 84:34) before obtaining total
ENA using TRIzol. cDNA library construction was made from
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                                                                                      organism="Trypanosoma
/mol_type="mRNA"
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1 (bases 1 to 337)

Whitfield,C.W., Band,M.R., Bonaldo,M.F., Kumar,C.G., Liu,L.,
Pardinas,J., Robertson,H.M., Soares,B. and Robinson,G.E.

Annotated expressed sequence tags and cDNA microarrays for studies
of brain and behavior in the honey bee
Genome Res. 12 (4), 555-566 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: generobi@life.uiuc.edu
This research was funded by the University of Illinois Critical
Thesearch Thitiatives Fund and a Burroughs-Wellcome Trust Innovation
Research Initiatives Fund and as Burroughs-Wellcome Trust Innovation
Award in Functional Genomics to G.E. Robinson and an NSF
Postdoctoral Fellowship in Bioinformatics to C.W. Whitfield.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Gene E. Robinson
Department of Entomology
University of Illinois
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Fax: 217 244 3499
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Apis mellifera
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BACKWARD: ATTAACCCTCACTAAAG
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Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apo
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/Glone lib="Bee Brain Normalized Library, BB16"
/note="Organ: brain; Vector: pT7T3-Pac; Site_1: EcoR1;
Site_2: Not1; The BB16 library was contributed by the
Soares laboratory and it was constructed and normalized
as described by Bonaldo, M.F., Lennon, G. and Soares,
M.B. (1996), Genome Research 6(9): 791-806. RNA was
prepared from dissected brains of adult worker bees of
                                                                                                                                                              /tissue_type="brain"
/dev_stage="adult worker honey bee"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                            organism="Apis mellifera"
/mol_type="mRNA"
/strain="mixed strains of
                                                                                                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                      sex="female"
                                                                                                                                                                                                                                                            clone="BB160004A20G12"
                                                                                                                                                                                                                                                                                                         .m. ligustica"
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory of Cellular and Developmental Biology NIDDK, National Institutes of Health 6 Center Drive MSC 2715, Bldg 6, Rm B1-13, Bethesda, MD 20892 USA Fax: (301) 496 5239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gene discovery using computational and microarray analysis of transcription in the drosophila melanogaster testis Genome Res. 10 (12), 2030-2043 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drosophila melanogaster (fruit fly)
Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 390)
Andrews,J., Bouffard,G.G., Cheadle,C., Lu,J., Becker,K.G. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AI945022

AI94502.y1 Drosophila melanogaster adult testis library Drosophila melanogaster adult testis library Drosophila melanogaster cDNA clone bs08b02 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequencing Center (NISC; see http://www.nisc.nih.gov).
Plate: 08 row: b column: 02
Seq primer: M13RP1 reverse primer (ABI).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://www.niddk.nih.gov/intram/people/boliver.htm Tissue isolation and library construction performed at the National Institute of Diabetes and Digestive and Kidney Diseases, NIH (see http://www.niddk.nih.gov/intram/people/boliver.htm). DNA sequencing and analyses performed by National Institutes of Health Intramural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         On Aug 17, 1999 this sequence version replaced gi:5735420
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AI945022.2 GI:9990370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: oliver@helix.nih.gov,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCGTCGAACGTTCGA 15
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/dev stage="1-5 day adult"
//dev stage="1-5 day adult"
//dev stage="1-5 day adult"
//dev stage="1-5 day adult"
//clone lib="Drosophila melanogaster adult testis library"
//note="Organ: testis; Vector: pBlueScript SK (Stratagene);
Site_1: EcoR I; Site_2: Xho I; Testes dissected from 1-5
day adult y(*) w(67cI)/y males raised at 25oC. RNA
isolated using Trizol (Life Technologies) and a single
round of Poly(A) + selection using Oligotex (Qiagen). cDNA
library constructed using Stratagene ZAP-cDNA syntesis
kit. Oligo dT-primed, size fractionated -1-6 kb, and
directionally cloned at EcoRI and XhoI in Uni-ZAP XR.
Following a single round of amplification pBlueScript SK
phagemids were mass excised. A distribution channel for
clones is being sought, but not currently available.
Requests for clones cannot be homored."
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1 59 c
                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /strain="y[*] w[67c1]/Y"
/db_xref="taxon:7227"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Drosophila melanogaster"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                clone="bs08b02"
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whitfield, C.W., Band, M.R., Bonaldo, M.F., Kumar, C.G., Liu, L.,

Pardinas, J., Robertson, H.M., Soares, B. and Robinson, G.B.

Annotated expressed sequence tags and cDNA microarrays for studies

of brain and behavior in the honey bee

Genome Res. 12 (4), 555-566 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BI510669
432 bp. mRNA linear EST 08-APR-2002
BB160003A20G01.5 Bee Brain Normalized Library, BB16 Apis mellifera
cDNA clone BB160003A20G01 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FORWARD: TAATACGACTCACTATAGGG
BACKWARD: ATTAACCCTCACTAAAG
Plate: BB160003A20 row: G column: |
Seg primer: AGCGGATAACAATTTCACACAGA
High quality sequence stop: 432
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Tel: 217 265 0309
Fax: 217 244 3499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Gene E. Robinson
Department of Entomology
University of Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apis mellifera (honeybee)
Apis mellifera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: generobi@life.uiuc.edu
This research was funded by the University of Illinois Critical
Research Initiatives Fund and a Burroughs-Wellcome Trust Innovation
Award in Functional Genomics to G.E. Robinson and an NSF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eŭkaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apoidea;
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         ø
/clone lib="Bee Brain Normalized Library, BB16"
/note="Organ: brain; Vector: pT773-Pac; Site 1: EccR1;
Site 2: Not1; The BB16 library was contributed by the
Soares laboratory and it was constructed and normalized
as described by Bonaldo, M.F., Lennon, G. and Soares,
M.B. (1996), Genome Research 6(9): 791-806. RNA was
prepared from dissected brains of adult worker bees of
various ages and various behavioral groups. "
a 69 c 102 g 98 t
                                                                                                                                                                                                                                                                                                                                               /organism="Apis mellifera"
/mol_type="mRNA"
/strain="mixed strains of European bees,
A.m. ligustica"
                                                                                                                                                                                                        /tissue_type="brain"
/dev_stage="adult worker honey bee"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                               /sex="female"
                                                                                                                                                                                                                                                                                      /clone="BB160003A20G01"
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71.4%;

Score 15;

DB 12;

Length 432;

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RESULT 6
BH816458
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AUTHORS
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                                                                                                                                      15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Mahairas GG, Wallace JC, Hood
High Throughput Sequencing Center
University of Washington
401 Queen Anne Avenue North, Seattle,
Tel: (206) 616-3618
Fax: (206) 616-3887
Email: jwallace@u_washington.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AQ623639
AQ623639
HS_5377_A2_F05_SP6E RPCI-11 Human Male BAC Library Homo sapiens genomic clone Plate=953 Col=10 Row=K, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seq primer: SP6
Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clones are derived from the human BAC library RPCI-11. For BAC library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm) or from Resear h Genetics (info@resgen.com). BAC end Web Server: http://www.htsc.washington.edu
Plate: 953 row: K column: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence-tagged connectors: A sequence approach to mapping
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1 (bases 1 to 609)
Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T.,
Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              scanning the human genome
Proc. Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                  TCGTCGAACGTTCGA 349
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                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop: 609.
Location/Qualifiers
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                                                                                                                                    Conservative
                                                                                                                                                                                                                 /clone_lib="RPCI-11 Human Male BAC Library"
/note="Vector: pBACe3.6; Site_1: EcoRI; Site_2: EcoRI;
Male blood DNA was isolated from one randomly chosen do
and partially digested with a combination of EcoRI and
EcoRI Methylase. Size selected DNA was cloned into the
pBACe3.6 vector at EcoRI sites"
124 c 162 g 125 t 13 others
                                                                                                                                                                                                                                                                                                                                                                       /mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                         /sex≃"male"
                                                                                                                                                                                                                                                                                                                                                         /clone="Plate=953 Col=10 Row=K"
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100.0%;
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84;
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1 (bases 1 to 842)

Loftus, B., Wang, Z., Roncaglia, P., Van Aken, S. a.

Gene discovery in the Entamoeba invadens genome
                                                                                                                                                                                           EINCQ64TR EI_10_12 KB Entamoeba genomic survey sequence.
BZ391656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tomkins, J.P., Luo, M., Hunt, G., Main, D., Frisch, D., Page, P.E., Guzman Nova, E. and Wing, R.A.

Development of Genomic Resources for honey bee (Apis mellifera L.):

BAC Library Construction, Preliminary STC Analysis, and

Identification of Clones Associated With Behavioral Traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AM Ba0021J24f Apis mellifera Apis mellifera genomic clone AM Ba0021J24f, genomic survey sequence. BH816458
                                                                                             Entamoeba invadens
Entamoeba invadens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: jtmkns@clemson.edu
Total hg bases = 231
Total primer: TAATACGACTCACTATAGGG
Seq primer: TAATACGACTCACTATAGGG
Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Apis mellifera

Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
                                                                          Eukaryota; Entamoebidae; Entamoeba.
                                                                                                                                                                     BZ391656.1 GI:30238193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 864 656 7288
Fax: 864 656 4293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              http://www.genome.clemson.edu/projects/stc/bee/AM_Ba/ To order clones from this library see http://www.genome.clemson.edu/orders " 134 c 156 g 207 t 4 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Apis mellifera"
/note="Vector: pCUGIBAC-1; Site_1: HindIII; Site_2: NotI;
For more details on library preparation and sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   analysis see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="Africanized honey
/db_xref="taxon:7460"
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'mol_type="genomic DNA"
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lab_host="E. coli"
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                           and Fraser, C.
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                                                                                                                                                                                                           Laboratory of Molecular Plant Biology
Stazione Zoologica 'Anton Dohrn'
Villa Comunale, I-80121, Napoli, Italy
                                                                                                                                                                                                                                                                                                               1 (bases 1 to 889)
Scala, S., Carels, N., Falciatore, A., Chiusano, M.L. and Bowler, C. Genome properties of the diatom Phaeodactylum tricornutum Plant Physiol. 129 (3), 993-1002 (2002)
                                                                                                                                                                             Villa Comunale, I-80121, I
Tel: 39 081 583 3268/3211
Fax: 39 081 764 1355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PTMM00709 Phaeodactylum tricornutum Uni-Zap XR Phaeodactylum tricornutum cDNA 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Eukaryotic Genomics
TIGR
9712 Medical Center Drive, Rockville, MD 20850,
                                                                                                                        Diatom EST Database(http://avesthagen.sznbowler.com)
Seq primer: T3 backward
                                                                                                                                                           Email: chris@alpha.szn.it
                                                                                                                                                                                                                                                                   Contact: Bowler C
                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; stramenopiles; Bacillariophyta; Bacillariophyceae; Bacillariophycidae; Naviculales; Phaeodactylaceae; Phaeodacty
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD375545.1 GI:31251159
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Class: sheared ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: enta@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 301-838-3543
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Other_GSSs: EINCQ64TF
Contact: Brendan Loftus
                                                                                                                                                                                                                                                                                                                                                                                                                                             Phaeodactylum tricornutum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /Glone lib="EI 10 12 KB"
/Glone lib="EI 10 12 KB"
/note="Vector: pHOS2; Site_1: BstXI; Total genomic DNA was solated from early log phase trophozoites of E. invadens IP-1 using a Qiagen plant DNA extraction kit. A shotgun medium-size plasmid library (average insert size of 10 - 12 kb) was generated by random mechanical shearing E. invadens genomic DNA, repairing the ends of DNA fragments with T4 Polymerase, adding BstXI adaptors and ligating into the BstXI site of a pUC-derived vector pHOS2."

a 152 c 140 g 269 t
/organism="Phaeodactylum tricornutum"
/mol_type="mRNA"
/db xref="taxon:2850"
/cell_line="CCMP632"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Entamoeba invadens"
/mol_type="genomic DNA"
/strain="IP-1"
                                                                                       ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'clone="EINCQ64"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  db_xref="taxon:33085"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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100.0%; Pred. No.
Live 0; Mismatcl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dq 688
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87;
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                                RESULT 10
CC235774/c
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TITLE
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BX455352/c
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   DEFINITION
                     POCOR
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                                                                                                                                                                          Best Local
Matches 1
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Best Local
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900 GAACGTTCGAGATGA 886
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CC235774 1220 bp
CH261-139L19_RM1.2 CH261 Gallus
                                                                                                                                                                                                     l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cgi-bin/cluster.cgi?seq=CSOBAK038CA04NM2&cluster=10667.f. Contact Feng Liang Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com/ InvitroGen Corporation 1600 Faraday Avenue Genoscope sequence ID: CSOBAK038CA04NM2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: seqref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen. This sequence belongs to sequence cluster 10667.f For
more information about this cluster, see
http://www.genoscope.cns.fr/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BX455352 Homo sapiens FETAL BRAIN Homo sapiens
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BX455352.1 GI:31019187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CSODF022YA13 3-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                GAACGTTCGAGATGA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 913)
                                                                                                                                                                                                                                                                                                    304
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                                                                                                                                                                                                 71.4%; Score 15; llarity 100.0%; Pred. No. Conservative 0; Mismatci
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a 247 c
                                                                                                                                                                                                                                                                                              /note="Organ: brain; Vector: pCMVSPORT_6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and ECORV sites of the pCMVSPORT 6 vector. Library was not normalized."

199 c 163 g 246 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="CSODF022YA13"
/rissue_type="FETAL BRAIN"
/dev_stage="fetal"
/clone_lib="Homo sapiens FETAL BRAIN"
/clone_lib="Homo sapiens FETAL BRAIN"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Phaeodactylum tricornutum Uni-Zap
/note="Vector: Uni-Zap XR vector; Site_1: Eco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Homo sapiens"
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Primates;
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Pred. No.
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                                                                                                                                                                                                 Pred. No. 88;
Mismatches 0;
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  gallus
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88;
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  linear
genomic clo
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    GSS 12-MAY-2003
e CH261-139L19,
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clone
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RI; Site_2:
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VERSION
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Kremitzki C., Higginbotham, J., V
Warren, W., Graves, T., Mardis, E.
Gallus gallus BAC End Reads
Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genome Sequencing Center
Washington University School of Medicine
Email: submissions@watson.wustl.edu
Insert Length: 182000 Std Error: 0.00
Seq primer: RM1 TACGACTCACTATAGGGAGA
Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence start: 473 High quality sequence stop: 541. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genomic survey sequence CC235774
                                                                                               Submitted (06-JAN-2003) Genoscope
BP 191 91006 EVRY cedex - FRANCE
                                                                                                                                                                                                                                 Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                       CNS09N54 108 bp mRNA linear HTC 08-JAN-2003 Single read from an extremity of a full-length cDNA clone made from Anopheles gambiae total adult females. 5-PRIME end of clone FKOAAC5AH09 of strain 6-9 of Anopheles gambiae (African malaria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Richard K. Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gallus gallus
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                                                                                                                                                                          Anopheles.
1 (bases 1 to 108)
                                                                                                                                                                                                                                                                                                  HTC
                                                                                                                                                                                                                                                                                                                                                         mosquito).
                                                                                                                                                                                                                   Neoptera; Endopterygota;
                                                                                                                                                                                                                                                                                                                  BX066068.1 GI:27639349
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                                                                                                                                        Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AACGTTCGAGATGAT 21
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                                                                             www.genoscope.cns.fr)
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/note="Vector: pTARBAC2.1; Site 1: EcoRI; Site_2: EcoRI;
/note="Vector: pTARBAC2.1; Site 1: EcoRI; Site_2: EcoRI;
CH261 Female Chicken library - For library and clone
ordering information: http://www.chori.org/bacpac"
a 330 c 171 g 399 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Gallus gallus"
/mol_type="genomic DNA"
/strain="Red Jungle Fowl"
/db_xref="taxon:9031"
/mol_type="mRNA"
                      organism="Anopheles gambiae"/
                                                          Location/Qualifiers
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100.0%; Fi
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=_lib="CH261"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Score 15; DB
%; Pred. No. 91;
0; Mismatches
                                                                                                                                                                                                                   Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wylie, K., Carter, J., McPherson, J., and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 29; Length 1220;
                                                                                               e - Centre National de Sequencage (E-mail : seqref@genoscope.cns.f)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 314 286 1455

Fax: 314 286 1810

Email: mmarra@watson.wustl.edu

Email: mmarra@watson.wustl.edu

Email: mmarra@watson.wustl.edu

Email: mmarra@watson.wustl.edu

Email: mmarra@watson.wustl.edu

PCR F: TGTAAAACGACCAGTGAGCCATTATGAGTATTCTTCCAAGGGTA

FOR B: CAGGAAACAGCTATGACCCTTATGAGTATTCTTCCAAGGGTA

SOUTCE: Washington University Genome Sequencing Center

PCR amplified DNA is available from Washington University Genome

Sequencing Center. Aliquots of the library may be requested from

Sumabara (pekemarc-lnb.cam.ac.uk).
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Hillier, L., Chiapelli, B., Chissoe, S., Clark, N., Couch, J., Dubuque, T., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Kuwabara, P., Le, M., Mardis, E., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Tan, M., Mardis, E., Waterston, R., Wohldmann, P. and Wilson, R., Fr., Trevaskis, E., Waterston, R., Wohldmann, P. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caenorhabditis briggsae
Caenorhabditis briggsae
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/db_xref="taxon:7165"
/clone="rkohAC54H09"
/plasmid="pME18S-FL"
/note="end : 5-pRIME"
a 31 c 34 g
                                                                                        /clone lib="Kuwabara Mixed stage C. briggsae"
/note="Vector: Lambda gt10; Site 1: EcoRI; Site 2: EcoRI;
Stage:mixed, Sex:hermaphrodite. Library construction:
First strand oligo(dT) primed. Second strand was as per
Gubler/Hoffman. Ligated to EcoRI adaptors. Library is
non-directional. Library is non-normalized. Library
constructed by P.B. Kuwabara. Additional details on
constructed by P.B. Kuwabara and S. Shah, NAR 22: 4414 - 4418 (1994). Adaptor
sequence: GAATTC CGTTGCTGTGTG
a 42 g 38 t
                                                                                                                                                                                                                                                                                                                                                         /organism="Caenorhabditis briggsae"
/mol_type="mRNA"
/strain="Gl6 Gujarat"
/db_xref="taxon:6238"
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66.7%; Score 14; DB 14; 100.0%; Pred. No. 2.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecular Biology Unit
International Livestock Research
P.O. Box 30709, Nairobi, Kenya
Tel: 254-2 630743
Fax: 254-2 631499
                                                                                                                                                                                 mosquito).
BX070096
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Djikeng,A., Donelson,J.E. and Majiwa,P.A.O.
Generation of expressed sequence tags as physical landmarks in the
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                                                                   Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                     Anopheles gambiae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
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                                 (bases 1 to 348)
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primer: T3 primer
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Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note=Tvector: Lambda ZAP II (Stratagene); Site 1: EcoxI; Site 2: XhoI; The mRNA was purified from a cloned population of bloodstream trypanosomes reexpressing the MVAT4 metacyclic variant surface glycoprotein (VSG). A unidirectional oligo dT-primed EcoRI/XhoI cDNA library was constructed in lambda ZAP II (Stratagene)."
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/mol_type="mRNA"
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_xref="taxon:31286"
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                                                                                                                                                                                                                                                                                                                                                                                                              Department of Entomology
University of Illinois
505 S. Goodwin Ave., Urbana, IL
Tel: 217 265 0309
Fax: 217 244 3499
Email: generobi@life.uiuc.edu
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Whitfield, C.W., Band, M.R., Bonaldo, M.F., Kumar, C.G., Liu, L.,
Pardinas, J., Robertson, H.M., Soares, B. and Robinson, G.E.
Annotated expressed sequence tags and cDNA microarrays for studies
of brain and behavior in the honey bee
Genome Res. 12 (4), 555-566 (2002)
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                                                                                                                                                                                                                                   Plate: BB170008B10 row: E column: Seq primer: AGCGGATAACAATTTCACACAGGA
                                                                                                                                                                                                                                                                                                                                   This research was funded by the University of Illinois Critical Research Initiatives Fund and a Burroughs-Wellcome Trust Innovation Award in Functional Genomics to G.E. Robinson and an NSF Postdoctoral Fellowship in Bioinformatics to C.W. Whitfield.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Apis mellifera (honeybee)
Apis mellifera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BB170008B10E04.5 Bee mellifera cDNA clone
                                                                                                                                                                                               High quality sequence stop: 387.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Gene E. Robinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apoidea;
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BACKWARD: ATTAACCCTCACTAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Web : www.genoscope.cns.fr)
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/clone="FKOAAC7ABO1"
/plasmid="pME18s-FL"
/note="end : 5-pRIME"
a 101 c 114 g
                                                                                               /organiam="Apis mellifera"
/mol_type="mRNA"
/strain="mixed strains of European bees,
/m. ligustica"
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/mol_type="mRNA"
/strain="6-9"
tissue_type="brain"
dev_stage="adult worker honey bee"
                                                            'db_xref="taxon:7460"
'clone="BB170008B10E04"
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                                          sex="female"
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Pred. No.
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3e+02;
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ORIGIN
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MEDLINE
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Best Local (
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414 bp mRNA linear EST 08-APR-2002
BB170012B20C02.5 Bee Brain Normalized/Subtracted Library, BB17 Apis
mellifera cDNA clone BB170012B20C02 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Gene E. Robinson
Department of Entomology
University of Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Apis mellifera (honeybee)
Apis mellifera
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apo
                                                                                                                                                                                                                                                                                  Plate: BB170012B20 row: C column: 02
Seg primer: AGCGGATAACAATTTCACACAGGA
High quality sequence stop: 414.
                                                                                                                                                                                                                                                                                                                                                       FORWARD: TAATACGACTCACTATAGGG
BACKWARD: ATTAACCCTCACTAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                       This research was funded by the University of Illinois Critical Research Initiatives Fund and a Burroughs-Wellcome Trust Innovation Award in Functional Genomics to G.E. Robinson and an NSF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pardinas,J., Robertson,H.M., Soares,B. and Robinson,G.E. Annotated expressed sequence tags and cDNA microarrays for studies of brain and behavior in the honey bee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 414)
Whitfield, C.W., Band, M.R., B.
Pardinas, J., Robertson, H.M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 217 265 0309
Fax: 217 244 3499
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                                                                                                                                                                                                                                                                                                                                                                                                             PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                              Postdoctoral Fellowship in Bioinformatics to C.W. Whitfield.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: generobi@life.uiuc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apidae,
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/note="Organ: brain; Vector: pT773-Pac; Site 1: EcoR1;
Site 2: Not1; This BB17 cDNA library was generated by
subtraction of the BB16 library with 4000 previously
sequenced clones. The BB16 library was contributed by the
Soares laboratory and it was constructed and normalized
as described by Bonaldo, M.F., Lennon, G. and Soares,
M.B. (1996), Genome Research 6(9): 791-806. RNA was
prepared from dissected brains of adult worker bees of
various ages and various behavioral groups. "
36 t
                                                                                                                                  /mol_type="mRNA"
/strain="mixed strains of European bees, predominantly
/strain="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="DH10B"
/tissue_type="brain"
/dev_stage="adult worker honey bee"
/lab_host="DH10B"
                                                                                               clone="BB170012B20C02"
                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                       sex="female"
                                                                                                                                                                                                                organism="Apis mellifera"
                                                                                                                   _xref="taxon:7460"
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; Pred. No. 3e+02;
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VERSION
                BASE COUNT
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ORGANISM
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CB333691 424 bp mRNA linear E
TgESTzyf67b10.y1 TgME49 B7 Tachyzoite cDNA Library 2
gondii cDNA clone TgESTzyf67b10.y1 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: toxo@watson.wustl.edu
Contact David Sibley (toxoest@borcim.wustl.edu) for further
information relating to organism, libraries, or clone availability.
Seq primer: -40RP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Toxoplasma EST Project
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sarcocystidae; Toxoplasma.
1 (bases 1 to 424)
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Bukaryota; Alveolata; Apicomplexa; Coccidia; Eimeriida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
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                93
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/dev_stage="Tachyzoite"
/lab_host="S.coli XL10"
/lab_host="S.coli XL10"
/clone_lib="TgME49 B7 Tachyzoite cDNA Library 2"
/clone_lib="TgME49 B7 Tachyzoite to the like the like
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/note="Organ: brain; Vector: pT773-Pac; Site_1: EcoR1;
Site_2: Not1; This BB17 cDNA library was generated by
subtraction of the BB16 library with 4000 previously
sequenced clones. The BB16 library was contributed by the
Soares laboratory and it was constructed and normalized
as described by Bonaldo, M.F., Lennon, G. and Soares,
M.B. (1996), Genome Research 6(9): 791-806. RNA was
prepared from dissected brains of adult worker bees of
various ages and various behavioral groups. "
a 100 c 104 g 110 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Toxoplasma gondii"
/mol_type="mRNA"
/db_xref="taxon:5811"
/clone="TgESTzyf67b10.y1"
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Pred. No. 3.1e+02;
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: HoltRA@celera.com
Plate: NU01004AZR row: K
Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1. (bases 1 to 433)
Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., R., Collins,F.H., Venter,J.C. and Hoffman,S.L. Celera Anopheles gambiae EST project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Celera Genomics
45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
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                                                                                                                                                                                                                                                                                  /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult morguitoes (mixed sex) frozen on liquid nitrogen.cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
a 109 c 137 g 76 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome) "
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="Adult"
/lab_host="DH10b"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /strain="RSP-ST (Reduced susc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone="19600449650107"
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A.Gam.ad.cDNA1 Anopheles
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3.1e+02;
thes 0;
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gambiae cDNA
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A clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                    113
Bode,H., Blumberg,B., Steele,R., Wigge,P., Gee,L., Nguyen,Q., Martinez,D., Kibler,D., Hampson,S., Clifton,S., Pape,D., Marra,M., Hillier,L., Martin,J., Wylle,T., Dante,M., Theising,B., Bowers,Y., Gibbons,M., Ritter,E., Bennett,J., Ronko,I., Tasgareishvili,R., Maguire,L., Kennedy,S., Waterston,R. and Wilson,R. Washire,L., Kennedy,S., Waterston,R. and Wilson,R. Contact: Hans Bode Contact: Hans Bode Washington University School of Medicine Washington University School of Medicine Washington University School of Medicine
                                                                                                                                                                                                                                      Hydra magnipapillata
Hydra magnipapillata
                                                                                                                                                                                                                                                                                                                                     CA302582 466 by taa13b12.x1 Hydra cDNA library
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1 (bases 1 to 439)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Mahairas,G.S., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. a:
                                                                                                                                                                                                          Eukaryota; Metazoa; Cnidaria;
Hydridae; Hydra.
                                                                                                                                                                                                                                                                          EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: jwallace@u.washington.edu
Sequence Tagged Connector
Plate: 2262 row: O column: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Mahairas GG, Wallace JC,
High Throughput Sequencing Center
University of Washington
401 Queen Anne Avenue North, Seatt
                                                                                                                                                                                                                                                                                          CA302582.1
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Location/Qualifiers
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Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: (206) 616-3618
Fax: (206) 616-3887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence-tagged connectors: A sequence
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                                                                                                                                                                                            (bases 1 to 466)
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Natl. Acad. Sci. U.S.A. 96 (17), 9739-9744 (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
/mol_type="genomic DNA"
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100.0%; Pr
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Pred. No. 3.1e+02;
0; Mismatches 0;
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CNS09GNZ

Single read from an extremity of a full-length cDNA clone made from Anopheles gambiae total adult females. 5-PRIME end of clone FKOAAC37DA02 of strain 6-9 of Anopheles gambiae (African malaria
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on obtaining a clone please contact: Hans Bode
Seq primer: Primer name ambiguous
High quality sequence stop: 426.
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Email: estGwatson.wustl.edu
Library was constructed by H. Bode and B. Blumberg DNA sequencing
Library was constructed by H. Bode and B. Center For information
                                                                                                                                                                                                                                                                                                                                                                                                                             Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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Location/Qualifiers
                                                                                                                    /db_xref="taxon:7165"
/clome="FKOAAC3"DA02"
/plasmid="pME18S-FL"
/note="end : 5-PRIME"
a 126 c 157 g
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/db xref="taxon:6085"
/db xref="taxon:6085"
/lab host="bH5alpha"
/clone lib="Hydra cDNA library"
/note="Vector: pSPORT6; Site 1: NotI; Site 2: SalI;
/inote="Vector: pSPORT6; Site 1: NotI; Site 2: SalI;
/ibraries prepared by Phil Wigge, Plant Molecular and
Cellular Biology, Lab (Detlev Weigel); The Salk Institute
for Biological Science, 10010 North Torrey Pines Road, La
Jolla, CA 92037."
                                                                                                                                                                                                               /organism="Anopheles
/mol_type="mRNA"
/strain="6-9"
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006 EVRY cedex - FRANCE
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(E-mail : seqref@genoscope.cns.fr
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Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Nematocera; Culio
                                                                                                                                                                                                                                                      8
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WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Ld
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 484)
Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuo, Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA09703.rl Stratagene mouse Tcell 937311 Mus musculus cDNA clon IMAGE:550608 5' similar to gb:X69392 60S RIBOSOMAL PROTEIN L26 (HUMAN); gb:X80699 M.musculus L26 mRNA (MOUSE);, mRNA sequence.
                                                                                                           17000687561542
19600449644818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: -28ml3 rev1 ET from Amersham
High quality sequence stop: 7.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                               BM636144.1
                                                                                               BM636144
                                                                                                                                           BM636144
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ilarity 100.0%;
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db_xref="taxon:10090"
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                                                                                                         Celera Genomics
45 w. Gude Dr., Rockville, MD 20850, USA
Tel: 2404533151
Fax: 2404534580
Email: HoltRA@celera.com
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Tel: 2404533151
Fax: 2404534580
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                                                                                                                                                                                                                                  1 (bases 1 to 492)
Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A.,
R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
Celera Anopheles gambiae EST project
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                                                                       Plate: NU01004ABV row: K
Seq primer: M13 Reverse.
                                                                                                                                                                                                     Contact: Holt R.A.
                                                                                                                                                                                                                      Unpublished
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/clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sall; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
/organism="Anopheles gambiae"
/mol_type="mRNA"
                                                      Location/Qualifiers
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/clone="19600449644818"
/dev_stage="Adult"
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/strain="RSP-ST (Reduced susc.
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45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 493)
Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., Kraft,C.L., Charlab,R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
Celera Anopheles gambiae EST project
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Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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17000687313490 A.Gam.ad.cDNA1 Anopheles gambiae cDNA
19600449671469 5', mRNA sequence.
BM644091
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Location/Qualifiers
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                                                             /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3', Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."

[www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                           /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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cDNA inserts >500 bp cloned directionally into pSport 1.

Not 1 site is 3'. Clones available through the Malaria

Research and Reference Reagent Resource Center
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/note="Vector: pSport1; Site_1: Sall; Site_2:
adult mosquitoes (mixed sex) frozen on liquid
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134 c
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lab_host="DH10b"
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'clone="19600449634767"
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tive 0; Mismatches
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BM611518/c
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neomrara: Endopterygota; Diptera; Nematocera; Culicoidea;
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Anopheles gambiae (African malaria mosquito)
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Eukaryota; Metazoa; Arthropoda; Hexapoda; In
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45 w. Gude Dr., Rockville,
Tel: 2404533151
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19600449683979 5', mRNA
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen 24 hours after human blood feeding. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)"
a 131 c 159 g 88 t
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/mol_type="mRNA"
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A.Gam.ad.cDNA.bloodl Anopheles
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  Arthropoda; Hexapoda; Insecta; Pterygota;
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1 (bases 1 to 504)

1 (bases 1 to 504)

Holt,R.A., Lin,J.-J., Murphy,S.D., Ev
Holt,R.A., Lin,J.-J., Murphy,S.D., Ev
Holt,R.A., Collins,F.H., Venter,J.C. and Ho
Holt,R.A., Collins,F.H., Venter,J.C. and Holt,R.A., Collins,F.H., Collin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM866297 512 bp mRNA linea mgcs005xE24f.b Magnaporthe grisea CS Uni-Zap XR grisea cDNA clone mgcs005xE24 5', mRNA sequence
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45 w. Gude Dr., Rockville,
Tel: 2404533151
                                                                                          Peterson Bldg, MS2132, College
Tel: 979 845 4831
Fax: 979 845 6483
                                                                                                                                                                                                               On Mar 7, 2002 this
Contact: Ebbole DJ
Email: d-ebbole@tamu.edu
Chromatogram file of this sequence is available, see contact person
;Best nr hit (April. 22, 2003) gb|EAA28529.1| hypothetical protein
[Neurospora crassa] 107 9e-49
                                                                                                                                                                     Department of Plant Pathology & Microbiology
Texas A&M University
                                                                                                                                                                                                                                                                                                                                             Ebbole, D.J., Yuan, J., Thomas, T.L., K. and Dean, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Magnaporthe grisea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: HoltRA@celera.com Plate: NU01004A9Y row: B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                   Sordariomycetes incertae sedis; Magnaporthaceae;
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/strain="RSP-ST (
chromosome)"
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/note="Vector: pSport1; Site_1: Sall
adult mosquitoes (mixed sex) frozen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="19600449722493"
/dev_stage="Adult"
/lab_host="DH10b"
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                                                                                                                                                                                                                                                                                                                      tags from the rice blast fungus,
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Hoffman, S.L
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Magnaporthe

SU

EST

06-MAY-2003 Magnaporthe

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Gaps

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Not1; Whole

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REFERENCE
AUTHORS
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FORWARD: T3 primer
BACKWARD: T7 primer
Plate: mgcs005 row:
Seg primer: T3.
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                                                                                                                                                                                                                                                                                                Contact: Ruan Y

Contact: Ruan Y

Laboratory of Molecular Biotechnology

Genome Institute of Singapore

Genome Park Road, The Capricorn #05-01, Singapore 117528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CB358730 11 ap mRNA linear EST 17-MAR-20
ZF001-P00024-DPE-F-A G12 GISZF001 Danio rerio cDNA clone
IMAGE:6900672 5' similar to fc24912.y1 Zebrafish WashU MPIMG EST
Danio rerio cDNA clone IMAGE:3722374 5', mRNA sequence.
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Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Danio.
                                                                                           Plate: ZF001-P00024-DPE-F-A
Seq primer: CCGCATAACTTGTATAGCA
                                                                                                                                              FORWARD: M13
BACKWARD: M13
                                                                                                                                                                                            PCR PRimers
                                                                                                                                                                                                              Email: gisry@nus.edu.sg
GIS Clone ID: ZF001-P00024-PP_M23
                                                                                                                                                                                                                                                                    Tel: +65 6827 5200
Fax: +65 6827 5201
                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mathavan, S., Wei, C., Thoreau, H., Chia, J.M. Genome Institute of Singapore, Zebrafish E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Danio rerio
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                                                                       quality sequence stop:
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/db_xref="taxon:148305"
/db_xref="taxon:148305"
/clone="mgsg005xE24"
/clone="tayon="tayon"
/clone=lib="Magnaporthe grisea CS Uni-Zap XR Library"
/clone=lib="Magnaporthe grisea CS Interest and primer and predominantly 5' reads T7 primer on XhoI side of insert.
Conidial library. Point incoulation of Guyll at center of oatmeal agar plate. Conidia were harvested after two weeks of growth. Sequences were processed by one of two methods.
of growth. Sequences were processed by one of two methods.
Where a full-length alignment to the M. grisea genome sequence was trimed according to the alignment, otherwise sequence quality was assessed using phredPhrap version 991019 and trimmed according to phd files (0.05) and for vector seqs."

18 a 161 c 119 g 114 t
                                           Location/Qualifiers
/organism="Danio rerio"
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/strain="Guy11"
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                          BP 44, Avenue Fontaule,
Tel: (33)468887309
Fax: (33)468887398
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Derelle, E., Ferraz, C., Lagoda, P., Eychenie, S., Cooke, R., F.
Sabau, X., Courties, C., Delseny, M., Demaille, J., Picard, A.
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Eukaryota; Viridiplantae; Chlorophyta; Prasinophyceae; Mamiellales;
Mamiellaceae; Ostreococcus.
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J. Phycol.
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                                                                                                                                                                                                                                                                                      Email: h.moreau@obs-banyuls.fr
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Moreau
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3' cloning site: Sfi B site GGCCGCCTCGGCC;
3' linker/adaptor sequence: same as the priming sequence; Average insert size: 2bb; For PCR insert analysis: Use M13 Forward and reverse primers; Library Amplified Recombinants (Inserts): 98%; Library complexity: 5x106; Full-length construction (method): SMART, a Clontech method; Library constructed by: S. Mathavan, Chia-Lin Wei, and Yijun Ruan Genome Institute of
/db_xref="taxon:70448" /clone="tcta23a09.b" /clone="tcta23a09.b" /clone="tcta23a09.b" /clone="0strecocccus tauri genomic shotgun library" /clone="Vector: Bluescript; Site_1: EcoRV; Site_2: EcoRV; Shotgun library prepared after Sonication of the genomic
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/note="Vector: pDNR-LIB, Site 1: Sfi A (GGCCATTACGGCC);
/note="Vector: pDNR-LIB, Site 1: Sfi A (GGCCATTACGGCC);
Site_2: Sfi B (GGCCGCCTCGGCC); Priming method: Sfi-(dT)30
Primed ; Priming sequence: 5.ATTCTACA GGCCGAGGCCGGCC
GACATG(T)30VN; Directionally cloned,
Sfi A site GGCCATTACGGCC;
Sfi A site GGCCATTACGGCC;
Sfi A site GGCCATTACGGCC;
                                                                                                                  /organism="Ostreococcus tauri"
/mol_type="genomic DNA"
/strain="OTTH0595"
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/dev stage="7 Different embryonic Stages( From just
fertilized Embryos to 72 hours just hatched baby fish)"
/lab_host="DH10B"
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/db_xref="taxon:7955"
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100.0%; Pr/
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Pred. No. 3.2e+02;
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; 3' cloning site: Sfi B
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BM618261/c
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Anopheles gambiae (African Mararre, ______
Anopheles gambiae
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Nematocera; Culicoidea;
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19600449662307 5', mRNA
BM618536
BM618536.1 GI:18916954
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BM618261
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Location/Qualifiers
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1: HoltRA@celera.com
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146 c 214 g 77 t
                                                                                                                                                                    /clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sall; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
138 c 174 g 90 t
                                                                                                                                                                                                                                                                                                                           chromosome) "
                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/strain="RSP-ST (Reduced
                                                                                                                                                                                                                                                                                     dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                             organism="Anopheles gambiae"
                                                                                                                                                                                                                                                                                                    clone="19600449662307"
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Gude Dr., Rockville,
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DEFINITION
                                                                                 Anopheles.

1 (bases 1 to 528)

Holt,R.A., Lin,J.-J., Murphy,S.D.,
,R., Collins,F.H., Venter,J.C. and
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Plate: NU01004NAK row: J
Seq primer: M13 Reverse.
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Holt,R.A., Lin,J.-J., Murphy,S.D.,
R., Collins,F.H., Venter,J.C. and
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19600449620529
Contact: Holt R.A.
Celera Genomics
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Tel: 2404533151
Fax: 2404534580
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/mol type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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lab_host="DH10b"
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'clone="19600449620529"
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Pred. No.
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. 3.2e+02;
                                                                                 Evans, C.A.,
Hoffman, S.L.
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                                                                                                        Kraft, C.L., Charlab
                                                                                                                                                                                                                                                                                                                                  cDNA clone
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MD 20850,

USA

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REFERENCE
AUTHORS
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VERSION
KEYWORDS
SOURCE
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TA99D10P
LOCUS
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ORIGIN
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JOURNAL
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                                                                                                                                                                                      Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTAt 10.1) was mechanically shear to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ហ
                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fax: 2404534580
Email: HoltRA@celera.com
Plate: NU01004HMK row:
Seq primer: M13 Reverse.
                                                                                                        Email: nelasyed@tigr.org
Details of T. brucei sequencing at the Sanger Centre
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hc Melville,S.E., Rajandream,M.A. and Barrell,B.G.

    brucei sheared genomic DNA
genomic survey sequence.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AL460276.1 GI:11862154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/dev_stage="Adult"
/lab_host="DH10b"
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/mol_type="genomic DNA".
/strain="TREU927"
/db_xref="taxon:5691"
                                                                 organism="Trypanosoma brucei"
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/strain="RSP-ST (Reduced susc.
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Pred. No. 3.2e+02;
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RESULT 36
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Matches 14; Conserv
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Best Local Similarity 100.0%;
                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
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                                                                                                                      336
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AQ407920 534 bp DNA linear GSS 17-MAR-
HS_5055_B2_E08_T7A RPCI-11 Human Male BAC Library Homo sapiens
genomic clone Plate=631 Col=16 Row=J, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Celera Genomics
45 w. Gude Dr., Rockville, MD 20850, USA
Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anopheles.

1 (bases 1 to 533)

Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., R., Collins,F.H., Venter,J.C. and Hoffman,S.L. Celera Anopheles gambiae EST project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 TCGTCGAACGTTCG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: HoltRA@celera.com Plate: NU01004N8E row: O Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished Contact: Holt R.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anopheles gambiae (African malaria mosquito)
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                                                                                                                      CGAACGTTCGAGAT 323
                                                                                                                                                      CGAACGTTCGAGAT 18
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                                                                                                                                                                                                                                                               /clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3', Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."

(www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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( 177 c 1
                                                                                                                                                                                                                                                                                                                                                                                                             /clone="19600449617956"
/dev_stage="Adult"
/lab_host="DH10b"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              db_xref="taxon:7165"
                                                                                                                                                                                     66.7%; Score 14; DB 12; 1 100.0%; Pred. No. 3.2e+02; ive 0; Mismatches 0;
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A.Gam.ad.cDNA1 Anopheles
5', mRNA sequence.
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Pred. No.
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. 3.2e+02;
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                                                                                                                                                                                                                     Length 533;
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                                                                                                                                                                                  ACCESSION
                                                                                                                                                                                                                       DEFINITION
                                                                                                                                                                                                                                                           AL695492/c
                                                                                                                                                                                                                                                                                RESULT 37
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Best Local S
Matches 14
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PUBMED
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AQ407920.1
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://www.htsc.washington.edu
Plate: 631 row: J column: 16
Seq primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: jwallace@u.washington.edu
Clones are derived from the human BAC library RPCI-11. For BAC
library availability, please contact Pieter de Jong
(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm)
or from Resear h Genetics (info@resgen.com). BAC end Web Server:
                                                          Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                          sequence.
AL695492
AL695492.1 GI:19615134
                                                                                                                                                                                                                 AL695492 NAP1 Anopheles gambiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 534.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High Throughput Sequencing Center University of Washington 401 Queen Anne Avenue North, Seattle, Tel: (206) 616-3618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence-tagged connectors: A sequence approach to mapping and scanning the human genome Proc. Natl. Acad. Sci. U.S.A. 96 (17), 9739-9744 (1999)
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Keller, A., Shaker, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 534)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Christophides, G.K.,
                                                                                                                                            EST.
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                                          Anopheles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Mahairas GG, Wallace JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hood, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      domo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                   AACGTTCGAGATGA 20
                        (bases 1 to 541)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (206) 616-3887
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="RPCI-11 Human Male BAC Library"
/note="Vector: pBACe3.6; Site_1: EcoRI; Site_2: EcoRI
Male blood DNA was isolated from one randomly chosen and partially digested with a combination of EcoRI an EcoRI Methylase. Size selected DNA was cloned into th pBACe3.6 vector at EcoRI sites"

a 71 c 149 g 135.t 6 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="Plate=631 Col=16 Row=J"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1. .534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                      66.7%;
    Blass, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                Score 14; DB 28; Le
s; Pred. No. 3.2e+02;
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    Zdobnov, E.M., Carmouche, R., Benes, V.
                                                                                                                                                                                                                     cDNA clone NAP1-P31-H-11-5,
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Young,J., Zhao,S.,
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Adams,M.D. and
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SOURCE
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VERSION
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BI511012/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                   AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              405
Email: generobi@life.uiuc.edu
This research was funded by the University of Illinois Critical
Research Initiatives Fund and a Burroughs-Wellcome Trust Innovation
Award in Functional Genomics to G.E. Robinson and an NSF
Postdoctoral Fellowship in Bioinformatics to C.W. Whitfield.
PCR PRimers
                                                                                                                                                                                                                                                                            1 (bases 1 to 544)
whitfield, C.W., Band, M.R., Bonaldo, M.F., Ku
Pardinas, J., Robertson, H.M., Soares, B. and
Annotated expressed sequence tags and cDNA
of brain and behavior in the honey bee
Genome Res. 12 (4), 555-566 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61511012 544 bp mRNA linear BB160004A20E05.5 Bee Brain Normalized Library, BB16 cDNA clone BB160004A20E05 5', mRNA sequence.
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European Molecular Biology Laboratory
Meyerhofstrasse 1, 69117 Heidelberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and Kafatos, F.C.
Anopheles gambiae EST, Eur
Unpublished
Contact: Christophides GK
                                                                                                                    505 S. Goodwin Ave., Urbana, Tel: 217 265 0309
Fax: 217 244 3499
                                                                                                                                                                              Contact: Gene E. Robinson
Department of Entomology
University of Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apis mellifera (honeybee)
Apis mellifera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: +49 6221 387-440
Fax: +49 6221 387-306
                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BI511012.1 GI:15361386
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: christop@embl-heidelberg
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ilarity 100.0%;
Conservative (
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/(clone lib="NAP1"
/(clone lib="NAP1"
/note="Vector: pT7T3D-Pac (Pharmacia); Site_1: NotI;
/note="Vector: pT7T3D-Pac (Pharmacia); Site_1: NotI;
/site_2: EcoRI; ESTs sequenced from the T7 priming site
that reads from the 5' end of cDNA. The NAP1 is a
directionally cloned and normalized, oligo-T primed cDNA
library constructed from a mixture of Anopheles gambiae
developmental stages according to: Bonaldo, Lennon &
Soares (1996): Normalization and Subtraction: Two
Approaches To Facilitate Gene Discovery, Genome Research
6, 791-806."
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/mol_type="mRNA"
/db_xref="taxon:7165"
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o. 3.2e+02;
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                                                                                                                                                                USA
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Apis mellifera
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REFERENCE
AUTHORS
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VERSION
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BM649585/c
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Best Local Similarity
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                                                                                                                  source
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                                                                                                                                                                                                                                                                                                           Anopheles.

Anopheles.

1 (bases 1 to 546)

1 (bases 1 to 546)

Holt,R.A., Lin,J.-J., Murphy,S.D., Ev
Holt,R.A., Lin,J.-J., Murphy,S.D., Ev
Holt,R.A., Collins,F.H., Venter,J.C. and Ho
,R., Collins,F.H., Venter,J.C. and Ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BM649585 546 bp mRNA linear EST 26 17000687369108 A.Gam.ad.cDNAl Anopheles gambiae cDNA clone 19600449617798 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                    Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plate: BB160004A20 row: E column: Seq primer: AGCGGATAACAATTTCACACAGGA
                                                                                                                                                                                                               Fax: 2404534580
                                                                                                                                                                                                                                                                                      Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BM649585.1 GI:18949096
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                                                                                                                                                                                                                                                                  Celera Genomics
                                                                                                                                                                                                                                                                                                         Unpublished
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                                                                                                                                                                                                                              w. Gude Dr., Rockville,
L: 2404533151
                                                                                                                                                 il: HoltRA@celera.com
te: NU01004NBE row: H
primer: M13 Reverse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Bee Brain Normalized Library, BB16"
/note="Organ: brain; Vector: pT773-Pac; Site_1: EcoR1;
Site_2: Not1; The BB16 library was contributed by the
Soares laboratory and it was constructed and normalized
as described by Bonaldo, M.F., Lennon, G. and Soares,
M.B. (1996), Genome Research 6(9): 791-806. RNA was
prepared from dissected brains of adult worker bees of
various ages and various behavioral groups."
                                     chromosome) "
                                                       /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc
                       db xref="taxon:7165"
                                                                                                                                  ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  strain="mixed strains of European bees, predominantly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
clone="19600449617798"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Apis mellifera"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lone="BB160004A20E05"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    xref="taxon:7460"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        stage="adult worker honey bee"
host="DH10B"
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Pred. No.
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                                                                                                                                                                       column:
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3.2e+02;
hes 0;
                                                         susc.
                                                                                                                                                                                                                                                                                                                                           Evans, C.A., Kraft, C.L., Charlab Hoffman, S.L.
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                                                         to Permethrin -
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                                                         std.
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KEYWORDS
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BM650226/c
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ORIGIN
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JOURNAL
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Best Local
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Holt,R.A., Lin,J.-J., Murphy,S.D.,
Holt,R.A., F.H., Venter,J.C. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     549 bp mRNA linear i
17000687370552 A.Gam.ad.cDNA1 Anopheles gambiae cDNA
19600449640068 5', mRNA sequence.
BM650226
                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: HoltRA@celera.com
Plate: NU01004AY4 row: I
Seq primer: M13 Reverse
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Tel: 2404533151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fax: 2404534580
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CGAACGTTCGAGAT 273
                              CGAACGITCGAGAT 18
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                                                                                                                                         /clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/note="Vector: pSport1; Site 1: Sall; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria
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                                                                                                                                                                                                                                                                                                                                              chromosome) "
                                                                                                                                                                                                                                                                               /clone="19600449640068"
/dev_stage="Adult"
/lab_host="DH10b"
                                                                                                                                                                                                                                                                                                                                                            'mol_type="mRNA"
'strain="RSP-ST (Reduced susc.
                                                                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                             organism="Anopheles gambiae"
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100.0%; Pr
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                                                             Score 14; DB 12;
Pred. No. 3.2e+02;
0; Mismatches 0;
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Pred. No. 3.2
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Evans, C.A.,
                                                                                                                                                                                                                                                                                                                                                                to Permethrin -
                                                                                            Length 549;
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RESULT 42
BM633500/c
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AUTHORS
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BM626834/c
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VERSION
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                                            REFERENCE
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JOURNAL
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Best Local :
                             AUTHORS
                                                                                                       ORGANISM
                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                         302
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17000687496213 A.Gam.ad.cDNA1
19600449642604 5', mRNA sequer
                                                                                                                                                                           BM633500 554 bp mRNA
17000687507362 A.Gam.ad.cDNA1 Anopheles
19600449648215 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                       v
         Anopheles.

1 (bases 1 to 554)

Holt.R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., Kraft,C.L., R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
                                                                     Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                     EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: HoltRA@celera.com Plate: NU01003CY3 row: B Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45 w. Gude Dr., Rockville, Tel: 2404533151 Fax: 2404534580
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BM626834.1 GI:18926345
                                                                                                                                                   BM633500.1
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Neoptera; Endopterygota; Diptera; Nematocera; Culio
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                                                                                                                                                                                                                                                                                                                      CGAACGTTCGAGAT 18
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Anopheles gambiae EST project
                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="mRNA"
/strain="RSP-ST (Reduced susc. to chromosome)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:7165"
/clone="19600449642604"
/dev_stage="Adult"
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                                                                                                                                                   GI:18933011
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Pred. No.
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3.2e+02;
hes 0;
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gambiae cDNA
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AUTHORS
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BH899413/c
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Best Local
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COMMENT
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                                                                                                                                                         BP 44, Avenue Fontaule,
Tel: (33)468887309
Fax: (33)468887398
                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 556)
Derelle, E., Ferraz, C., Lagoda, P., Eychenie, S., Cooke, R., Rega
Sabau, X., Courties, C., Delseny, M., Demaille, J., Picard, A. and
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Ots00/480 Ostreococcus tauri genomic
tauri genomic clone otb06b09.b 5', g
BH899413
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Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                        Ostreococcus tauri
Eukaryota; Viridiplantae; Chlorophyta; Prasinophyceae; Mamiellales;
                                                                                                                                                                                                                                     Contact: Moreau H
Laboratoire Arago
                                                                                                                                                                                                                                                                         J. Phycol.
                                                                                                                                                                                                                                                                                                           DNA libraries for sequencing the genome of Ostreococcus tauri (Chlorophytae, Prasinophyceae): the smallest free-living euka
                                                                                                                                                                                                                                                                                                                                                                                                                         Mamiellaceae; Ostreococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ostreococcus tauri
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Seq primer: M13 Reverse
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Celera Genomics
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                                                                                                                                            Email: h.moreau@obs-banyuls.fr
                                                                                                                                                                                                                      CNRS UMR 7628
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                                                                                                                           primer: forward
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/note="Vector: pSport1; Site 1: Sall; Site 2: Not1; Whole adult mosquices (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
               /organism="Ostreococcus
/mol_type="genomic DNA"
/strain="OTTH0595"
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/strain="RSP-ST (Reduced
                                                                                     ocation/Qualifiers
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lab_host="DH10b"
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                                                                       . .556
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                                                                                                                                                                                                                                                                           38 (6),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GI:22550908
_xref="taxon:70448"
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3.2e+02;
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Determination of clone end sequences from Trypanosoma brucei GUTat 10.1 sheared DNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Clones are derived from the Trypanosoma brucei GUTat 10.1 sheared DNA library constructed at TIGR. Clones will be available for distribution through Research Genetics, Alabama, USA. Sheared DNA end sequences search page: http://www.tigr.org/tdb/mdb/tbdb/. Seg_primer: M13-Forward
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Sheared DNA-85E2.TF Sheared DNA Trypanosoma
Sheared DNA-85E2, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Other_GSSs: Sheared DNA-85E2.TR
Contact: Najib M. El-Sayed
Department of Eukaryotic Genomics
The Institute for Genomic Research
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Trypanosoma brucei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: nelsayed@tigr.org
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                                       /clone_lib="Sheared DNA"
/note="Vector: pUC18; Site_1: Smal; Constructed at The
/note="Vector: pUC18; Site_1: General File File
Institute for Genomic Research (TIGR), Rockville, MD.
Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically
sheared to give a tight size distribution (approx 2 kb).
The v + i method used for the library construction is
described in detail in Smith, H.O. and Venter, J.C.
(Making small insert libraries for whole genome shotgun
sequencing projects. In Genome Sequencing: A Practical
Approach, eds. M. Vaudin and B. Barell, Oxford University
Press, 1999)."
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/note="Vector: Bluescript; Site 1: EccRV; Site 2: EccRV;
Shotgun library prepared after sonication of the genomic
DNA. Blunt ligation in EccRV site of Bluescript. Size
selection of the inserts after agarose electrophoresis
between 1 and 3 kb."
                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:5691"
/clone="Sheared DNA-85E2"
                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="TREU927/4 GUTat 10.1"
                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Trypanosoma brucei"
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3.2e+02;
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RESULT 46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               561 bp mRNA linear EST 26.
17000687504190 A.Gam.ad.cDNA1 Anopheles gambiae cDNA clone
19600449668153 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Celera Genomics
45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                          BM600003
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Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
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                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: HoltRA@celera.com
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                                                                                                                                                                                                                                                                                    /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."

187 g 95 t
                                                                                                                                                                                                                                                                                                                                                                                                            /clone="19600449668153"
/dev_stage="Adult"
/lab_host="DH10b"
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/strain="RSP-ST (Reduced susc.
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                 GI:18898107
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48 A.Gam.ad.cDNA.blood1 A
58 5', mRNA sequence
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100.0%; Pred. No. 3.:
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                                                          Anopheles gambiae
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Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A.,
R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
Celera Anopheles gambiae EST project
                                                                                                                                                                                                                                                                                                                           BM621639
BM621639.1 GI:18920057
EST.
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Plate: NU01004AYG row: K
Seq primer: M13 Reverse.
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Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Endopterygota; Diptera; Nematocera; Culicoidea;
     Plate: NU01004AYL row: I Seq primer: M13 Reverse.
                                                                           45 w. Gude Dr., Rockville, Tel: 2404533151
                                                                                                                                                                             1 (bases 1 to 569)
Holt,R.A., Lin,J.-J.,
,R., Collins,F.H., Ver
                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Nematocera; Culio
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Anopheles gambiae
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19600449637012 5', mRNA
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                                                                                                            Celera Genomics
                                                                                                                               Contact: Holt R.A.
                                                                                                                                                              Celera Anopheles gambiae
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                                                         2404534580
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                                      HoltRA@celera.com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=Tvector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen 24 hours after human blood feeding. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)" a 153 c 183 g 94 t
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/dev_stage="Adult"
/lab_host="DH10b"
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/mol_type="mRNA"
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Pred. No.
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. 3.2e+02;
ches 0;
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Culicoidea;
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AUTHORS
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BM619639/c
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KEYWORDS
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JOURNAL
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Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A.,
,R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
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17000687440685 A.Gam.ad.cDNA1 Anopheles gambiae cDNA clone
19600449657860 5', mRNA sequence.
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Plate: NU01004AB6 row: J
Seq primer: M13 Reverse.
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45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
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/clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
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chromosome)"
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                                                                                                                                                /clone="19600449657860"
/dev_stage="Adult"
/lab_host="DH10b"
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'strain="RSP-ST (Reduced
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lab_host="DH10b"
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Pred. No. 3.2e+02;
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EST 25-FEB-2002

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REFERENCE
AUTHORS
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JOURNAL
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A1667035 577 bp
fc24912.y1 Zebrafish WashU MPIMG
IMAGE:3722374 5', mRNA sequence.
A1667035 A1667035.1 GI:4805391
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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Plate: NU01004187 row: E
Seq primer: M13 Reverse.
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Tel: 2404533151
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/note="Vector: pSport1; Site 1: Sall; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
151 c 188 g 96 t
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/mol_type="mRNA"
/strain="RSP-ST (Reduced susc. to Permethrin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone="19600449663794"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 stage="Adult"
                                                                                                                                                                                                                Score 14; DB 12; 1; Pred. No. 3.2e+02; 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clark, M., Johnson, S.L., Lehrach, H., Lee, R., Li, F., Marra, M., Eddy, S., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwoo, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: zbrafish@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
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4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished Contact: Stephen L. Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and Wilson, R.
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1 (bases 1 to 577)
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                                                                                                66.7%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                           double-stranded cDNA was ligated to Sal I adaptors (BRL), digested with Not I and cloned into the Not I and Sal I sites of the pSPORTI vector (BRL). Library was constructed by Matthew Clark (Lehrach lab; ICRP, London and Max Planck Institut fuer Molekulare Genetik, Berlin). cDNAs for EST analysis were selected following oligonucleotide hybridization fingerprinting of arrayed clones from zebrafish late somitogenesis (26 ss), adult liver or embryonic shield stage (5.6 h) libraries. Fingerprint data were used to computationally cluster cDNAs, and a single cDNA from each cluster was chosen for sequencing. In some cases multiple members of the same cluster were sequenced to assess clustering parameters or single clones were sequenced additional times to assess quality control."
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/lab_host="XL1-blue MRF"
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db_xref="taxon:7955"
clone="IMAGE:3722374"
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                                                                                                                                           Nippostrongylus brasiliensis
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida;
Trichostrongyloidea; Heligmonellidae; Nippostrongylinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., R., Collins,F.H., Venter,J.C. and Hoffman,S.L. Celera Anopheles gambiae EST project
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Unpublished
                                                            Harcus, Y. and Maizels, R.M.
                                                                                                                                                                                                                                                                                                               BM279238.1 GI:17972496
                               Sequence survey of Nippostrongylus
                                                                                                                        Nippostrongylus.
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te: NU01004AYT row: O
primer: M13 Reverse.
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Research and Reference Reagent (www.malaria.mr4.org)."
154 c 185 g 101 t
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/mol_type="mRNA"
/strain="RSP-ST (Reduced susc
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/dev_stage="Adult"
/lab_host="DH10b"
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REFERENCE
AUTHORS
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BF157792
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VERSION
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JOURNAL
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Institute of Cell, Animal a
University of Edinburgh
Ashworth Labs, West Mains R
Tel: +44 131 650 5511
Fax: +44 131 650 5450
                Email: zbrafish@watson.wustl.edu
Library constructed by Dr. Sumio Sugano and Dr. Koichi Kawakami DNA
Sequencing by: Washington University Genome Sequencing Center Clone
distribution information can be found through the I.M.A.G.E.
Consortium/LLNL, send email to: info@image.llnl.gov
Seq primer: T3 ET from Amersham
                                                                                                                                                                                                                                                                                           i (bases 1 to 583)
Clark, M., Johnson, S.L., Lehrach, H., Lee, R., Li, F., Marra, M., Eddy, S., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylle, T., Underwoo, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., K., Steptoe, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  583 bp mRNA line
fl44g06.yl Sugano Kawakami zebrafish DRA Danio
IMAGB:3816778 5', mRNA sequence.
BF157792
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1 (hasea 1 ro 52)
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Danio rerio
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Seq primer: Ml3
                                                                                                                                         Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                    Contact: Stephen L. Johnson Washington University School of Medicine
                                                                                                                                                                                                                                                              and Wilson,R.
WashU Zebrafish EST Project 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BF157792.1 GI:11052991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The library was prepared by Murray Selkirk, Imperial College, London. Sequencing was performed by Yvonne Harcus, Edinburgh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: r.maizels@ed.ac.uk
High quality sequence
                                                                                                                                                                               4444 Forest Park Parkway, Box 8501,
                                                                                                                                                                                                                                             Unpublished
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/clone="Nb adl 04E02"
/dev_stage="adult"
/clone_lib="Nippostrong
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mol_type="mRNA"
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Ostariophysi; Cypriniformes
                                                                                                                                                                                     Louis,
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RESULT 54
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Best Local Similarity
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                                                                                                                                                                                                                                                  Celera Genomics
45 w. Gude Dr., Rockville,
Tel: 2404533151
                                                                                                                                                                                                                                                                                                                                                                                  Anopheles.

1 (bases 1 to 588)

r.in,J.-J
                                                                                                                                                                                                                                                                                                                                                                                                                               Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     588 bp mRNA linear EST 26
17000687502159 A.Gam.ad.cDNA1 Anopheles gambiae cDNA clone
19600449617012 5', mRNA sequence.
                                                                                                                                                                                               Email: HoltRA@celera.com Plate: NU01004N8A row: H
                                                                                                                                                                                                                                                                                                            Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                               Unpublished
                                                                                                                                                                                                                                                                                                                                                               Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., Kraft,C.L., Charlab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM630963
                                                                                                                                                                                                                                       Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                              Celera Anopheles gambiae
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/lab_host="DH10B (phage resistant)"
/lab_host="DH10B (phage resistant)"
/clone_lib="Sugano Kawakami zebrafish DRA"
/note="Wector: pME18S-FL3; Site_l: DraIII (CACTGTGTG);
/note="Wector: pME18S-FL3; Site_l: DraIII (CACTGTGTG);
/note="Wector: pME18S-FL3; Site_l: DraIII (CACTGTGTG);
/note="Wector: pME18S-FL3; Site_l: Strand cDNA was primed with an oligo(dT) primer [ATGTGGCCTTTTTTTTTTTTTTT];
/double-stranded cDNA was ligated to a DraIII adaptor [TGTTGGCCTACTGG], digested and cloned into distinct DraIII sites of the pME18S-FL3 vector (5' site CACTGTGTG, 3' site CACCATGTG). XhoI should be used to isolate the cDNA insert. Size selection was performed to exclude fragments <1.5kb. Library constructed by Dr. Sumio Sugano (University of Tokyo Institute of Medical Science) and kindly donated by Dr. Xoichi Kawakami. Custom primers for sequencing: 5' end primer CTTCTGCTCTAAAAGCTGCG and 3' end primer CGACCTGCAGCTCGAGCACA. "

39 a 148 c 152 g 143 t 1 others
                                                                                                                                                                                                              HoltRA@celera.com
                                                                     /organism="Anopheles gambiae"
/mol type="mRNA"
/strain="RSP-ST (Reduced susc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="mixed (one male and unfertilized eggs)"
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/clone="19600449617012"
/dev_stage="Adult"
/lab_host="DH10b"
                                                        db_xref="taxon:7165"
                                                                                                                                                             ocation/Qualifiers
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/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone="IMAGE:3816778"
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; Pred. No. 3.20
0; Mismatches
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3.2e+02;
hes 0;
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                                                                                        to Permethrin -
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AUTHORS
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BM649878/c
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Matches 1
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                  CGAACGTTCGAGAT 429
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5 CGAACGTTCGAGAT 18
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1 (bases 1 to 592)

Holt,R.A., Lin,J.-J., Murphy,S.D.,
,R., Collins,F.H., Venter,J.C. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 592 bp mRNA
17000687369980 A.Gam.ad.cDNA1 Anopheles
19600449651232 5', mRNA sequence
EM649878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Celera Genomics
45 w. Gude Dr., Rockville,
Tel: 2404533151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: HoltRA@celera.com plate: NU010049W6 row: Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anopheles gambiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Celera Anopheles gambiae EST project
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                                                                                                                             /note=Tvector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. CDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."

158 c 199 g 97 t
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                                                                                                           chromosome) "
                                                                                                                                                                                                                                                                                             /clone="19600449651232"
/dev_stage="Adult"
/lab_host="DH10b"
                                                                                                                                                                                                                                                                                                                                                                                   mol_type="mRNA" strain="RSP-ST (Reduced susc. to Permethrin -
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                                                                                                                                                                                                                                                                          clone_lib="A.Gam.ad.cDNA1"
                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Anopheles gambiae"
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100.0%; Pr
                                                           100.0%;
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                                  Score 14; DB 12;
Pred. No. 3.2e+02;
0; Mismatches 0;
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Pred. No. 3.2
0; Mismatches
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3.2e+02;
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REFERENCE
AUTHORS
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BM622534/c
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BM624679/c
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Best Local
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17000687491666 A.Gam.ad
19600449643241 5', mRNA
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Anopheles.

1 (bases 1 to 595)

Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., R., Collins,F.H., Venter,J.C. and Hoffman,S.L. Celera Anopheles gambiae EST project
                                                                              EST.
Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45 w. Gude Dr., Rockville, Tel: 2404533151
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BM624679.1 GI:18924190
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                                                                                                                                                                                     BM622534
                                                                                                                                                                                                   17000687476438 A.Gam.ad.
19600449643086 5', mRNA
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Seq primer: M13 Reverse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/strain="RSP-ST (Reduced susc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              db_xref="taxon:7165"
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Pred. No.
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.cDNA1 Anopheles gambiae
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3.2e+02;
hes 0;
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                                  Kraft, C.L.,
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JOURNAL COMMENT
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BM632552/c
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KEYWORDS
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   595 bp mRNA
17000697504447 A.Gam.ad.cDNA1 Anopheles
19600449625402 5', mRNA sequence.
BM632557
                                                                                                                                                                                                                                                                                               1 (bases 1 to 595)
Holt,R.A., Lin,J.-J., Murphy,S.D.,
,R., Collins,F.H., Venter,J.C. and
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Fax: 2404534580
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45 w. Gude Dr., Rockville,
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                                                                                                                                                                                45 w. Gude Dr.,
Tel: 2404533151
Fax: 2404534580
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                                                                                                                                              Email: HoltRA@celera.com Plate: NU01004NBW row:
                                                                                                                                                                                                                                 Celera Genomics
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                                                                                                                                                                                                                                                                  Unpublished
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/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
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/mol type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
                                /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc
chromosome)"
/db_xref="taxon:7165"
/clone="19600449625402"
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lab_host="DH10b"
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Pred. No.
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3.2e+02;
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Hoffman, S.L
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KEYWORDS
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BM639133/c
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Best Local S
Matches 14
                                                                                 Query Match
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431
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: HoltRA@celera.com
Plate: NU01004I8A row: I column: 17
Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Holt, R.A., Lin, J.-J., Murphy, S.D., Evans, C.A., R., Collins, F.H., Venter, J.C. and Hoffman, S.L. Celera Anopheles gambiae EST project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Bukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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EST.
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19600449666203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Holt R.A.
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CGAACGTTCGAGAT 418
                           CGAACGTTCGAGAT 18
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                                                                                                                                                                  144
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                                                                   66.7%; Score 14; DB ilarity 100.0%; Pred. No. 3.: Conservative 0; Mismatches
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/clone_lib="A.Gam.ad.cDNA1"
/clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
                                                                                                                                                      /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site 1: Sall; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
158 c 194 g 104 t
                                                                                                                                                                                                                                                                                                           /clone="19600449666203"
/dev_stage="Adult"
/lab_host="DH10b"
                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Anopheles gambiae"
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nes 14; Conserv
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BM621612
17000687447485 A.Gam.ad.cDNA1 Ar
19600449636865 5', mRNA sequence
BM621612
BM621612.1 GI:18920030
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: (206) 616-3618
Fax: (206) 616-3887
Email: jwallace@u.washington.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             http://www.htsc.washington.edu
Plate: 989 row: H column: 12
Seq primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clones are derived from the human BAC library RPCI-11. For BAC library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering bac.htm or from Resear h Genetics (info@resgen.com). BAC end Web Server:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High Throughput Sequencing Center
University of Washington
401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-3618
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601 bp DNA linear GSS 30-AUG HS_5413_B2_D06_T7A RPCI-11 Human Male BAC Library Homo sapiens genomic clone Plate=989 Col=12 Row=H, genomic survey sequence.
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1 (bases 1 to 601)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Mahairas,G.G., Wallace,J.C., Smith,K., Startzell,S., Adams,M.D. au
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Mahairas GG, Wallace JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10449764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            scanning the human genome
Proc. Natl. Acad. Sci. U.S.A. 96 (17), 9739-9744 (1999)
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                                                                                                                                                                                                                                                                                 66.7%;
ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: pBACe3.6; Site_1: EcoRI; Site_2: EcoRI; Male blood DNA was isolated from one randomly chosen donor and partially digested with a combination of EcoRI and EcoRI Methylase. Size selected DNA was cloned into the pBACe3.6 vector at EcoRI sites"

114 c 116 g 195 t 10 others
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone_lib="RPCI-11 Human Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Homo sapiens"
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Pred. No. 3.2
0; Mismatches
                                                            sequence
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. 3.2e+02;
ches 0;
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BH416973/c
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BH416973
BH416973.1 GI:17
GSS.
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                                                                                                                                                                                                               Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: HoltRA@celera.com Plate: NU01004AYL row: C Seq primer: M13 Reverse.
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Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
               Email: walbot@stanford.edu
Plate: 1007052 column: 30
                                                855 California Ave, Palo
Tel: 650 723 2227
Fax: 650 725 8221
                                                                                               Department of Biological Sciences
Stanford University
                                                                                                                                      Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                     1007052A03.x1 1007 - RescueMu Grid H Zea mays genomic, genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
                                                                                                                                                                 Maize genomic sequences found using engineered
                                                                                                                                                                                     Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGAACGTTCGAGAT 18
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transposon-tagged.
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162 c 201 g 88 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
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/lab_host="DH10b"
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3.2e+02;
hes 0;
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BM634523/c
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                                                                                                                                                                                                                                                                                                       Celera Genomics
45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                                                                                                                                                                                                                                                                                                                                                                   [bases 1 to 614]
Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A.,
.R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
Celera Anopheles gambiae EST project
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Anopheles gambiae
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EST.
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17000687509206 A.Gam.ad.cDNA1 Anopheles
1960049654085 5', mRNA sequence.
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                                                                                                                                                                                                                                              Plate: NU01004A9N row: P
Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                       Email: HoltRA@celera.com
                                                                                                                                                                                                                                                                                                                                                                                 Contact: Holt R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neoptera; Endopterygota;
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llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="1007 - RescueMu Grid H"
/clone_lib="1007 - RescueMu Grid H"
/note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site 1: BamHI; Site 2: BgIII;
RescueMu is a 4.9 kb, modified maize Mu transposon
designed to allow plasmid rescue from total genomic DNA.
Mu elements insert preferentially into transcription
units. For more information on RescueMu, go to the web
site 'www.zmdb.iastate.edu' and follow the links for
'RescueMu.' Grid H was grown at Berkeley in 2001. DNA
was extracted from leaf punches, double digested using
BamHI and BgIII, and ligated to form circular plasmids.
DH10B cells were transformed and then screened on LB
plates with ampicillin."
/clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sal1; Site_2:
/note="Vector: pSport1; Site_1: Sal1; Site_2:
adult mosquitoes (mixed sex) frozen on liquid
                                                                                                                               chromosome) "
                                                      /db xref="taxon:7165"
/clone="19600449654085"
/dev_stage="Adult"
/lab_host="DH10b"
                                                                                                                                                  /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /dev stage="adult"
/lab_host="DH10B"
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/cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
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hes 0;
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RESULT 64
BM625508/c
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RESULT 65
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Fax: 2404534580
Email: HoltRA@celera.com
Plate: NU0100418B row: O c
Seq primer: M13 Reverse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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17000687493212 A.Gam.ad.cDNA1 Ano
19600449662124 5', mRNA sequence.
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                                                                                                                                                                                                /clone_lib="A.Gam.ad.cDNA1"
/note="Vector: pSportl; Site_1: Sall; Site_2: Notl; Whole adult mosquitoes (mixed sex) Trozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                                                                                               /organism="Anopheles gambiae"
/mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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164 C 202 g 95 t
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/lab_host="DH10b"
                                                                                                                                                                                                                                                                                                                                    /clone="19600449662124"
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                                                                                                                               Score 14; DB 12; ; Pred. No. 3.2e+02;
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shes 0;
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Hoffman, S.L.
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                     TITLE
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                                                                                                                                                                                                                                                                                                                                                                                    510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-3618
Fax: (206) 616-3887
Email: jwallace@u.washington.edu
Clones may be purchased from Research Genetics (info@resgen.com)
BAC end Web Server: http://www.htsc.washington.edu
Plate: 4867 row: I column: 1
Seq primer: T7
                                                                                                            Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda, Hexapoda, Insecta, Pteryg
Neoptera; Endopterygota, Diptera, Nematocera, Culicoidea,
                                                                                                                                                                                                                                          BM591817 619 bp n
17000687435900 A.Gam.ad.cDNA.blood1
19600449730335 5', mRNA sequence.
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1 (bases 1 to 617)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T.,
Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AQ880074 617 bp DNA linear GSS 09-NOV-199: HS_4867_A1_E01_T7A CIT Approved Human Genomic Sperm Library D Homo sapiens genomic clone Plate=4867 Col=1 Row=I, genomic survey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Mahairas GG, Wallace JC,
High Throughput Sequencing Center
University of Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      scanning the human genome Proc. Natl. Acad. Sci. U.S.A. 96 (17), 9739-9744 (1999) 99380589
Celera Anopheles
Unpublished
                                   Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., Kraft,C.L., R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
                                                                                                                                                                                                           BM591817.1
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Location/Qualifiers
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                                                                                            Anopheles.
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ilarity 100.0%;
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/note="Organ: sperm; Vector: pBeloBAC11; BAC Clones in
E-Coli DH10B"
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/db_xref="taxon:9606"
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; Pred. No. 3.2e+02;
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CC120030/c
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                                                                                                                                                                                                                                                                 Other GSSs: NDL.76020.SP6
Contact: Brendan Loftus
Department of Eukaryotic Genomics
                                                                                                                                                                                                                                                                                                                            Aedes aegypti (yellow fever mosquito)
Aedes aegypti
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aedes.
1 (bases 1 to 627)
Loftus,B., Shetty,J., Knudson,D. and Severson,D.
BAC end sequencing of Aedes aegypti
                                                                                                                                                                                                                         Tel:
                                                                                                                                                                                                                                                                                                                                                                                                                                           GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC120030 627 bp DNA linear GSS NDL.76020.T7 Notre Dame Liverpool Aedes aegypti genomic NDL.76020, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: HoltRA@celera.com Plate: NU010049UV row: I Seq primer: M13 Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
                                                                                                                                            Seq primer: T7
Class: BAC ends.
                                                                                                                                                               Email: enta@tigr.org
Library was provided by
Seq primer: T7
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CC120030.1
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Celera Genomics
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301-838-0208
                                                                                                                                                                                                                                     Medical Center Drive,
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/clone lib="A.Gam.ad.cDNA.blood1"
/clone lib="A.Gam.ad.cDNA.blood1"
/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen 24 hours after human blood feeding. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)"
a 160 c 199 g 114 t
/db xref="taxon:7159"
/clone="NDL.76020"
/clone_lib="Notre_Dame_Liverpool"
/note="Vector: pECBAC1; Site_1: Hind
                                                            /mol_type="genomic [
/strain="Liverpool"
                                                                              /organism="Aedes aegypti"
/mol_type="genomic DNA"
                                                                                                                          Location/Qualifiers
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/dev_stage="Adult"
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/mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
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100.0%; Pred. No. 3.2e+02;
Mismatches 0;
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FORWARD: M13 Reverse (AGCGGATAACAATTTCACACAGGA)
BACKWARD: T7PL (CTCACTATAGGGCGAATTGG)
Seq primer: SAC (GGGAACAAAGCTGGAG).
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Institute of Cell, Animal and Population Biology
University of Edinburgh
Ashworth Labs, King's Buildings, West Mains Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: mark.blaxter@ed.ac.uk
The Trichuris muris EST dataset (including the TMC clustering information) is available on the www at http://www.ed.ac.uk/(tilde
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MBTmAdA198SAC Trichuris muris (parasitic r
Trichuris muris cDNA clone MBTmAdA198 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: +44 131 650 6760 Fax: +44 131 670 5450
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A survey of genes expressed in adults
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llarity 100.0%;
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                                                                                                   /note="Vector: Lambda Zap II; Site_1: EcoRI (5'end);
Site_2: Xho[3'end]; Trichuris muris is a nematode
parasite of rodents related to the human whipworm
Trichuris trichiura. The library was constructed from
Trichuris muris adults (Edinburgh 'E' strain) maintained
in mice, and was provided by Dr. Richard Grencis,
University of Manchester."

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by David Severson at the University of Notre Dame and
Hongbin Zhang"
148 c 149 g 171 t
                                                                                                                                                                                                                                                                             /dev_stage="adult"
/clone_lib="Trichuris muris (parasitic nematode) mixed
adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                         L. .628
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'db_xref="taxon:70415"
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                                                                                                                                                                                                                                                                                                                                        sex="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Trichuris muris"
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100.0%; Prr
Score 14; DB 9; ; Pred. No. 3.2e+ 0; Mismatches
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hes 0;
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nematode) mixed adult
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Best Local
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                                                                                                                                                                                                                                                    382
                                 Anopheles gambiae
Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda;
Eukaryota; Metazoa; Diptera; Nematoco
                                                                                                                                17000687475928 A.Gam.ad.cDNA1 Ar.
19600449654480 5', mRNA sequence
BM622163
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Anopheles.
1 (bases 1 to 629)
Holt, R.A., Lin, J.-J.,
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Email: HoltRA@celera.com
Plate: NU01004N8I row: N
Seq primer: M13 Reverse.
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Holt,R.A., Lin,J.-J., Murphy,S.D., Evans,C.A., R., Collins,F.H., Venter,J.C. and Hoffman,S.L.
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Anopheles gambiae
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
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BM641079.1 GI:18940590
EST.
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19600449621382 5', mRNA sequer
                                                                                    Anopheles gambiae (African malaria mosquito)
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                                                                                                                                                                                                                                                                                                                                                                      cDNA inserts >500 bp cloned directionally into psport 1.
Not 1 site is 3'. Clones available through the Malaria
Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
163 c 208 g 104 t
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/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab host="DH10b"
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/mol_type="mRNA"
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 Murphy, S.D.,
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. 3.2e+02;
:hes 0;
                                             Nematocera;
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Evans, C.A.,
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                                                       Insecta;
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                                           ecta; Pterygota;
Culicoidea;
Kraft, C.L., Charlab
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BM655453/c
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Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda, Hexapoda, Insecta,
Neoptera, Endopterygota, Diptera, Nematocera, Culio
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                                                                                                                                                                                                                                                         Unpublished
                                                                                                                                                                                                                                                                          Celera Anopheles gambiae EST project
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19600449627614 5', mRNA
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                                                                                                                            Email: HoltRA@celera.com Plate: NU010049VY row: A
                                                                                                                                                               Tel: 2404533151
Fax: 2404534580
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Fax: 2404534580
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45 w. Gude Dr.,
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Location/Qualifiers
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/mol_type="mRNA" /strain="RSP-ST (Reduced susc. chromosome)"
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/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."

[Www.malaria.mr4.org)."
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/mol_type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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/clone="19600449654480"
/dev_stage="Adult"
/lab_host="DH10b"
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Culicoidea;
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BM625565/c
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45 w. Gude Dr., Rockville,
Tel: 2404533151
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Holt,R.A., Lin,J.-J., Murp,R., Collins,F.H., Venter,
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17000687493286 A.Gam.ad.cDNA1 Anopheles gambiae cDNA clone
19600449629012 5', mRNA sequence.
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Plate: NU01004ABZ row: L
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult morguiroes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                 /organism="Anopheles gambiae"
/mol type="mRNA"
/strain="RSP-ST (Reduced susc.
chromosome)"
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Culicoidea;
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BM654222/c
                                                                             SOURCE
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17000687490421
19600449718154
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45 w. Gude Dr., Rockville,
Tel: 2404533151
Fax: 2404534580
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Holt,R.A., Lin,J.-J., Murphy,S.D., Ev.,R., Collins,F.H., Venter,J.C. and Hc Celera Anopheles gambiae EST project
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Anopheles gambiae
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea;
            Neoptera; Endopterygota; Anopheles.
                                                           Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
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19600449631529
                                            Eukaryota;
                                                                                                         BM597192.1
                                                                                                                          BM597192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: HoltRA@celera.com Plate: NU010049W1 row: D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BM654222.1 GI:18953733
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milarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                  /clone lib="A.Gam.ad.cDNA1"
/note="Vector: pSport1; Site_1: Sal1; Site_2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen. cDNA inserts >500 bp cloned directionally into pSport 1. Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center (www.malaria.mr4.org)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Anopheles gambiae"
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/strain="RSP-ST (Reduced susc.
chromosome)"
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lab_host="DH10b"
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'clone="19600449631529"
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                            Metazoa; Arthropoda; Hexapoda; Insecta; Pteryg
Endopterygota; Diptera; Nematocera; Culicoidea;
                                                                                                         GI:18895295
                                                                                                                           633 bp mk
21 A.Gam.ad.cDNA.blood1 A
34 5', mRNA sequence
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A.Gam.ad.cDNA1 Anopheles
5', mRNA sequence.
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. 3.2e+02;
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Hoffman, S.L.
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es gambiae cDNA clone
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45 w. Gude Dr., Rockville, MD
Tel: 2404533151
Fax: 2404534580
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                                                                                                                                                                                                                                                                                      Anopheles.

1 (bases 1 to 636)

1 (bases 1 to 636)

Holt, R.A., Lin, J.-J., Murphy, S.D., Evans, C.A., Kraft, C.L., Charlab, R., Collins, F.H., Venter, J.C. and Hoffman, S.L.
                                                                                                                                                                                                                                                                                                                                                                  Anopheles gambiae (African malaria mosquito)
Anopheles gambiae
Eukaryota, Metazoa; Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Endopterygota, Diptera, Nematocera, Culicoidea,
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17000668938404 A.Gam.ad.cDNA1 Anopheles gambiae cDNA clone
19600448419407 5', mRNA sequence.
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Plate: NU010049VS row: M
Seq primer: M13 Reverse.
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45 w. Gude Dr., Rockville,
Tel: 2404533151
                                                                                                                            Email: HoltRA@celera.com
Plate: NU010048I4 row: C
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/organism="Anopheles gambiae"
/mol_type="mtNA"
/strain="RSP-ST (Reduced susc. to Permethrin chromosome)"
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/mol_type="mRNA"
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CGAACGTTCGAGAT 356
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/note="Vector: pSport1; Site 1: Sal1; Site 2: Not1; Whole adult mosquitoes (mixed sex) frozen on liquid nitrogen.
cDNA inserts >500 bp cloned directionally into pSport 1.
Not 1 site is 3'. Clones available through the Malaria Research and Reference Reagent Resource Center
(www.malaria.mr4.org)."
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lab_host="DH10b"
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'clone="19600448419407"
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The present invention describes an immunomodulatory polynucleotide (I) comprising an immunostimulatory sequence (ISS). Also described: (I) an immunomodulatory composition comprising (I); (2) an immunomodulatory polynucleotide/microcarrier (IMP/MC) complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and (3) a kit comprising (I). (I) has antiallergic, antiasthmatic, virucide, antibacterial and protozoacide activities, and can be used as a modulator of immune response. (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune
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The present invention describes an immunomodulatory polynucleotide (I) comprising an immunostimulatory sequence (ISS). Also described: (I) an immunomodulatory composition comprising (I); (2) an immunomodulatory polynucleotide/microcarrier (IMP/MC) complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and (3) a kit comprising (I). (I) has antiallergic, antiasthmatic, virucide, antibacterial and protozoacide activities, and can be used as a modulator of immune response. (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune response, especially an allergy or asthma, or an infectious disease. (I)
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immunomodulatory composition comprising (1); (2) an immunomodulatory polynucleotide/microcarrier (IMP/MC) complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and (3) a kit comprising (I). (I) has antiallergic, antiasthmatic, virucide, antibacterial and protozoacide activities, and can be used as a modulator of immune response. (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune response, especially an allergy or asthma, or an infectious disease. (I) is also useful for increasing interferon-gamma (IFN-gamma) in an
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                                                                            The present invention describes an immunomodulatory polynucleotide (I) comprising an immunostimulatory sequence (ISS). Also described: (1) an immunomodulatory composition comprising (I)? (2) an immunomodulatory polynucleotide/microcarrier (IMP/MC) complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and (3) a kit comprising (I). (I) has antiallergic, antiasthmatic, virucide, antibacterial and protozoacide activities, and can be used as a modulator of immune response. (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune response, especially an allergy or asthma, or an infectious disease. (I)
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Pred. No
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The invention relates to a method for modulating cell death or reducing DNA damage in an eukaryotic cell by contacting the cell with an agent that modulates the biological activity of DNA-dependent protein kinase (DNA-PK). The invention also relates nucleic acids which modulate the immune response binding to Ku antigen, resulting in activation of DNA-PK. The method is useful for modulating cell death or reducing DNA damage in an eukaryotic cell, for treating any disorder resulting from a genotoxic insert to a cell e.g., necrosis, apoptosis. The method is also useful for treating cell death-related indications such as Alzheimer's disease,
                                                                                                                                                                                                                                                                                                                                                      Raz
                                                                                                                                                                                                                       Modulating cell death or reducing DNA damage in eukaryotic cells, useful for reducing cell death in individual or organ, comprises contacting cell with agent modulating biological activity of DNA-dependent protein kinase
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17-JAN-2001; 2001US-262321P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-MAY-2001; 2001WO-US14508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cell death; DNA damage; DNA-dependent protein kinase; DNA-PK; necrosis; immune response; apoptosis; Alzheimer's disease; Parkinson's disease; rheumatoid arthritis; inflammation; osteoporosis; myocardial infarction; liver disease; reperfusion injury; carcinoma; multiple sclerosis; stroke; amyotrophic lateral sclerosis; Acquired Immune Deficiency Syndrome; AIDS; head injury damage; aplastic anaemia; tumour; organ transplantation; cerebral infarction; follicular lymphomas; systemic lupus crythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Double-stranded immunostimulatory oligodeoxynucleotide (ISS-ODN).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD24905 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                     Example 1; Page 31; 57pp; English
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                                                                                                                                                                                                                                                                                                                    2002-062244/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               infection; glomerulonephritis; apoptosis;
s; immunostimulatory oligodeoxynucleotide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A; 4 C; 5 G; 5 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                      Takabayashi
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                                                                                                                                                                                                                                                                                                                                                      7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               autoimmune disorder; ISS-ODN; ds.
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Best Local S
Matches 15
                                                                                    (IMM),
'PN) that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infarction, bypass heart surgery, organ transplantation. The method is also useful for treating follicular lymphomas, carcinomas, autoimmune disorders (systemic lupus erythematosus), hormone dependent tumours, immune mediated glomerulonephritis; apoptosis and viral infections. The present sequence is immunostimulatory oligodeoxynucleotide (ISS-ODN) used for identifying ISS-binding protein, which is used in the exemplification of the invention
immune response against an antigen to a level greater that the host immune response to either the IMM, antigen or ISS-PN alone. These responses to ISS-PN/IMM conjugates are particularly acute during the important early phase of the host immune response to an antigen. The ISS-PN/IMM conjugates boost both humoral (antibody) and cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  central nervous system inflammation, osteoporosis, degenerative liver disease, cerebellar degeneration, reperfusion injury, multiple sclerosis, amyotrophic lateral sclerosis, myocardial infarction, head injury damage, acquired immunodeficiency syndrome (AIDS), aplastic anaemia, cerebral
                                                                                                                            This is the nucleotide sequence of DY1018, which is conjugated to beta-gal to form ISS-PN/IMM, comprising an immunomodulatory molecule
                                                                                                                                                                                                         New immunomodulatory compositions - comprising an antigen conjugated to a polynucleotide that contains an immunostimulatory sequence
                                                                                                                                                                                                                                                                                                                                                                                    09-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DY1018; beta-gal; ISS-PN/IMM; antigen; immune response; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-SEP-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parkinson's disease, rheumatoid arthritis, septic shock, sepsis, stroke, central nervous system inflammation, osteoporosis, degenerative liver
                                                                                                                                                                             Example 1; Page 36; 69pp; English.
                                                                                                                                                                                                                                                          WPI; 1998-261028/23.
                                                                                                                                                                                                                                                                                         Carson DA,
                                                                                                                                                                                                                                                                                                                                                       11-OCT-1996;
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                                                                                                               which comprises an antigen
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                                                                              which comprises an antigen conjugated to a polynucleotide sequence at contains at least one immunostimulatory nucleotide sequence. The conjugate synergistically boost the magnitude of the host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAACGTTCGAGATGA 20
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                                                                                                                                                                                                                                                                                                                        CALIFORNIA.
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                                                                                                                                                                                                                                                                                         Roman M;
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Pred. No.
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This is the ISS-ODN DY1018 nucleotide sequence.

The invention relates to a method for preventing or reducing antigen-stimulated, granulocyte-mediated tissue inflammation in a mammal, the proventing an immunostimulatory oligonucleotide (ISS-ODN), where:

CC (a) reduction in, or the absence of, a Th2 type immune response is comeasured; or (b) there is a reduction or absence of other clinical signs of inflammation in the host after antigen challenge. The method is used to reduce or suppress granulocyte-mediated inflammation in a host tissue, cand to modulate the host's immune responsiveness to an antigen, cand to modulate the subject suffers from asthma, nasal polyposis, callergic rhinitis, atopic dermatitis, allergic conjunctivitis, corect cutaneous basophil hypersensitivity. Unlike prior art treatment by candidated inmunisation, the method is an antigen-independent method, carries risk of anaphylaxis, and IL-5 which actually encourages granulocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                granulocyte-mediated tissue inflammation; Th2 type immune response; immune responsiveness modulation; idiopathic hypereosiniphilic syndrome; cutaneous basophil hypersensitivity; ISS-ODN; asthma; nasal polyposis; allergic chinitis; atopic dermatitis; allergic conjunctivitis;
                                                                                                                                                                                                                                                                                                                                      Reducing antigen-stimulated granulocyte-mediated inflammation
                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-312404/26
                                                                                                                                                                                                                                                                                                                                                                                                          Ray E;
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                                                                                                                                                                                                                                                                                                    Page 30; 69pp; English.
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100.0%; Pr
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Best Local :
                                                                                                                                                      sequences are selected from the group consisting of AACGTTCC, ACGTTCG, and GACGTTCG. The immunomodulatory sequences are used to treat patients needing immune regulation, such as those suffering from cancer, an allargic disease and asthma. They are also used to prevent infectious diseases such as influenza, herpes, hepatitis B, human immunodeficiency and papillomavirus, Hemophilus influenza, Mycobacterium tuberculosis and Bordetella pertussis, malarial plasmodia, Leishmania, Trypanosoma and Schistosoma. The immunomodulatory sequences are used to screen for human immunostimulatory activity by incubating macrophage cells and the oligonucleotide; and determining the relative amount of Thi-biased cytokines in the supernatant. Sequences AAV80104 to AAV80116 represent oligonucleotides that were tested for immunostimulatory activity. These
                                                               were used in experiments for the stimulation of cytokine production and were found to lack immunostimulatory activity. The invention provides specific claimed examples (AAV80096-103) of immunomodulatory sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 29; 63pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunostimulatory oligonucleotides regulate the immune system - and contain an immune-stimulating octanucleotide sequence; for treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunomodulatory; immunostimulatory;
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                        AAV80097 standard; DNA;
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influenza; herpes; M. tuberculosis; ss;
dia; Leishmania; Trypanosoma; Schistosom
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunostimulatory oligonucleotides regulate the immune system -contain an immune-stimulating octanucleotide sequence; for treat cancer, allergic and infectious diseases
Immunomodulatory; immunostimulatory; octanucleotide; immune regulatio ISS: cancer; allergy; asthma; hepatitis B infection; papillomavirus; Schiman immunodeficiency virus; influenza; herpes; M. tuberculosis; ss; B. pertussis; malaria; plasmodia; Leishmania; Trypanosoma; Schistosom
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                                                                                Immunomodulatory oligo comprising an ISS sequence.
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                                                                                                                                                                                    AAV80102
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Matches 15
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 15-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC (APC) with an intigen, comprising contacting antigen. Presenting cells (CAPC) with an immunostimulatory oligodeoxynucleotide (ISS-ODN). The APCs thus treated have enhanced antigen presenting capabilities compared to cartigen-activated APCs. APCs with enhanced antigen presentation compared to capabilities then present the antigen to T-cells. The method is useful compared to capabilities then present the antigen to T-cells. The method is useful compared to capabilities then present the ISS-ODN is used to enhance the tumour capabilities are inducing T-cell antigen presenting capacity of tumour cells. Thereby inducing T-cell cartivation, and is therefore useful for treating tumours. Additionally, common cells treated with an ISS-ODN ex vivo are useful as vaccines. CC ISS-ODN treated APCs are induced to take up antigen through upregulation of comportance co-stimulatory factors (B7 and CD40), to produce co-stimulatory factors (B7 and CD40), to complex (MHC) Class I and II expression and CC provide cell-to-cell adhesion through upregulation of intercellular contact of APC with antigen alone. The present sequence represents contact of APC with antigen alone. The present sequence represents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence of a stabilised oligonucleotide with antitumour activity
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                                                                                                                                                                           19-MAR-1999;
                                                                                                                                                                                                                              17-MAR-2000; 2000WO-FR00676
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunostimulatory oligonucleotide; tumour; anaplasia;
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Pred. No.
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19;
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RESULT 15
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Barackman i
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29-JUL-1999;
28-OCT-1999;
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                                                    emulsion which is
                                                      Microemulsion having an adsorbent surface comprising a microdiemulsion consisting of a metabolizable oil and an emulsifying which is a detergent, useful as a vaccine to treat bacterial,
                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                       human immunodeficiency virus; cytomegalovirus; CMV; in: rabies virus; cholera; diphtheria; tetanus; pertussis;
                                                                                                                                                                                                                                                                                                                                                                                                                                  CpG oligonucleotide; CpG motif; adjuvant; microdroplet emulsion; microemulsion; adsorbent microparticle; vaccine; Thi immune response; viral infection; hacterial infection; parasitic infection; HCV; HBV; hepatitis C virus; hepatitis B virus; herpes simplex virus; HSV; HIV;
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                                                                                                               WPI; 2000-587123/55
                                                                                                                                                                                                                                                                         09-FEB-2000; 2000WO-US03331.
                                                                                                                                                                                                                                                                                                                                  WO200050006-A2
                                                                                                                                                                                                                                                                                                                                                                                          Helicobacter pylori; Haemophilus influenzae; malaria; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CpG adjuvant
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                                        parasitic
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                                       consisting of a metabo
a detergent, useful as
sitic infection -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide,
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99US-0146391.
99US-0161997.
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Pred. No. 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                      influenza
                                                                                    microdroplet
                                                                                                                                                         Singh
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17;

Page 40; 95pp; English

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RESULT 16
AAA14467
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Best Local Similarity
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Immunostimulatory oligonucleotide adjuvant induces mucosal immunity an antigen in a mammalian host through production of secretory immunoglobulin {\tt A} -
                                                                                                                              WPI; 2000-303647/26.
                                                                                                                                                                                             Raz E,
                                                                                                                                                                                                                                                                                                                           05-OCT-1998;
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                                                                                                                                                                                                                                                         (REGC ) UNIV CALIFORNIA.
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The invention relates to a method of inducing mucosal immunity to

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Claim 3; Page 16; 65pp; English

Claim 8; Page 21; 64pp; English.

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RESULT 17
AAA38065
ID AAA38
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Matches
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                                                             Anti-HIV composition comprises immunostimulatory polynucleotides and HIV glycoprotein gp120 useful for modulating, stimulating an immune response against HIV in an HIV infected individual
                                                                                                                                                                                                 Tighe H,
                                                                                                                                                                                                                                                                                           09-OCT-1998;
07-OCT-1999;
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gp120; human immunodeficiency virus; HIV; immune response; infectic
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                                                                                                                                                         WPI; 2000-317846/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200021556-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                development;
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99US-0415186.
                                                                                                                                                                                                                                                                                                                                                               99WO-US23677.
                                                                                                                                                                                                   Schwartz D, Takabayashi
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0; Mismatches
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RESULT 18
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The present invention relates to an immunostimulatory composition comprising a human immunodeficiency virus (HIV) antigen, and an immunomodulatory polymucleotide comprising an immunostimulatory sequence (ISS). This sequence represents an ISS that can be used in the composition. An immunostimulatory composition which comprises a gp120 conjugated to an immunomodulatory polymucleotide, or is proximately associated to an immunomodulatory polymucleotide, or is proximately producing aspecific immune response against gp120 in an individual by producing anti-gp120 antibodies or gp120 specific cytotoxic T cells. It is also used for suppressing or delaying development of HIV infection in an individual infected with HIV or an individual at risk of infection
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                                                                                                                                                                                                                                                                                                                                                                                                 Anti-HIV composition comprises immunostimulatory polynucleotides and HIV glycoprotein gp120 useful for modulating, stimulating an immune response against HIV in an HIV infected individual
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07-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                             Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory sequence; ISS; immunomodulator; glycoprotein 120;
gp120; human immunodeficiency virus; HIV; immune response; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DYNA-) DYNAVAX TECHNOLOGIES CORP.
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                                                                                                                                                                                                                                                                                                                                        Page 16; 65pp;
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                                               conjugated to an immunomodulatory polynucleotide, or is proximately associated to it and not conjugated, is used for modulating or stimulating a specific immune response against gp120 in an individual by producing anti-gp120 antibodies or gp120 specific cytotoxic T cells. It is also used for suppressing or delaying development of HIV infection in an individual infected with HIV or an individual at risk of infection with HIV, respectively. It is also used for treating an individual infected with HIV in need of immune modulation.
                                                                                                                                                                                                                                                                    immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS). This sequence represents an ISS that can be used in the composition. An immunostimulatory composition which comprises a gp120
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gp120; human immunodeficiency virus; HIV; immune response; infectio
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                                                                                                                                                                                                                                                                                                                                                         comprising a human immunodeficiency virus (HIV) antigen,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 17; 65pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schwartz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "5-Bromocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISS) #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Takabayashi K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 21;
. 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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RESULT 20
AAA38072
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                                    conjugated to an immunomodulatory polynucleotide, or is proximately associated to it and not conjugated, is used for modulating or stimulating a specific immune response against gp120 in an individual by producing anti-gp120 antibodies or gp120 specific cytotoxic T cells. It is also used for suppressing or delaying development of HIV infection in an individual infected with HIV or an individual at risk of infection with HIV, respectively. It is also used for treating an individual
                                                                                                                            The present invention relates to an immunostimulatory composition comprising a human immunodeficiency virus (HIV) antigen, and an immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS). This sequence represents an ISS that can be used in the composition. An immunostimulatory composition which comprises a gp120
Sequence 22 BP; 6 A; 3 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                      Anti-HIV composition comprises immunostimulatory polynucleotides HIV glycoprotein gp120 useful for modulating, stimulating an immu response against HIV in an HIV infected individual -
                                                                                                                                                                                                                                                                                                                      Tighe
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07-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
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development;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory sequence; ISS; immunomodulator; glycoprotein 120;
gp120; human immunodeficiency virus; HIV; immune response; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA38072 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                (DYNA-) DYNAVAX TECHNOLOGIES CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 GAACGITCGAGATGA 20
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                          with HIV in need of immune modulation.
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                                                                                                                                                                                                                                                                                                                     Raz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                              Page 17; 65pp; English.
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99US-0415186.
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                                                                                                                                                                                                                                                                                                                     Schwartz D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             71.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "5-Bromocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "5-Bromocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15; ; Pred. No.
                                                                                                                                                                                                                                                                                                                      Takabayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Query Match

71.48;

Score 15;

DB 21;

Length 22;

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RESULT 21
AAZ55876
                                                                                        Sequences AAZ55876-255877 and AAZ55880-Z55886 represent immunomodulatory CC oligonucleotides comprising an immunostimulatory sequence (ISS, e.g., AACGTC, AACGTT, AGCGTT, AGCGTT, GACGTT, GACGTT, GACGTT, GACGTT, GACGTT, GACGTT, AGCGTT, AGCGTT, GACGTT, GACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                therapy, in treatment of allergy (including asthma), fo vigorous cellular response (against a virus, bacterium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 35; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel immunomodulatory oligonucleotide used
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01-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 response, e.g. to tumor antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9962923-A2
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Inh immune response; cytotoxic T-cell; cytokine; cancer; allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunomodulatory oligonucleotide SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DYNA-) DYNAVAX TECHNOLOGIES CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
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99US-0324191.
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9..16
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RESULT 22
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                                                                                                                                    polynuclectides which may be used to ameliorate inflammation of the gastrointestinal tract by administering a nucleic acid comprising one of these sequences. These polynucleotides all comprise an immunomodulatory nucleotide sequence of 5 °CpG-3 (I) The incleatides may be used for ameliorating or reducing gastrointestinal inflammation e.g. chronic or acute gastrointestinal inflammatory bowel disease; diarrhoea, rectal bleeding, weight loss; to reduce colon weight and colon lesions; to reduce a colonic inflammation. The immunomodulatory polynucleotides treat inflammatory bowel disease satisfactorily and effectively and have little or no toxicity even at a high dosage of 50000 micro-g. They also reduce the risk of
                                                                                                                                                                                                                                                                                               The
                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                        Claim 7;
                                                                                                                                                                                                                                                                                                                                               Ameliorating gastrointestinal inflammation e.g. inflammator disease involves administering an immunomodulatory nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                 Raz E,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ulcerative colitis; Crohn's disease; inflammatory bowel disease; diarrhoea; rectal bleeding; weight loss; colon; weight; lesion; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunomodulatory polynucleotide 1018.
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                                                                                                                                                                                                                                                                                               sequences given in AAH43338-48 represent immunomodulatory
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                                                                                                                            cancer
             GAACGTTCGAGATGA
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GAACGTTCGAGATGA 22
                                                                                                                                                                                                                                                                                                                        Page 28; 58pp; English.
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                                                 Conservative
                                                                                                 BP; 6 A; 3 C; 7 G; 6 T; 0 other;
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                                                                                                                           by treating ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation; gastrointestinal tract;
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                                                         71.4%;
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Pred. No.
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orv nucleic acid -
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RESULT 23
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                                                                                                                                         The invention is used to prevent and treat respiratory syncytial virus infection of the lower respiratory tract and other viruses including influenza virus, rhinovirus, adenovirus, measles virus, virus, parainfluenza virus, rubella virus, poxvirus, parvovirus, hantavirus and varicella virus. A kit for carrying out the administration is also included. Unlike the prior art antiviral agent ribavirin, which is a potential teratogen, the invention provides a treatment which does not carry unacceptable side effects. Other prior art medicaments treat the symptoms only, whilst the invention treats the infection. The present sequence is an ISS of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory sequence; ISS; ds; antiviral; immunogen; respiratory syncytial virus; RSV; influenza virus; rhinovirus; adenovirus; measles virus; mumps virus; parainfluenza virus; rubella virus; parvovirus; hantavirus; varicella virus.
                                                                                                                                                                                                                                                                                           The invention relates to suppressing a respiratory syncytial virus (RSV) infection in an exposed individual, comprising administering a polynucleotide comprising an immunostimulatory sequence (ISS) comprising the sequence (ISS) where an RSV antigen is not administered.
                                                                                                                                                                                                                                                                                                                                                                                                 Suppressing a respiratory syncytial virus infection by administering immunostimulatory sequence at the site of infection is useful to prevent and treat lower respiratory tract viral infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-MAR-2000;
09-MAR-2001;
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                                                                       Similarity
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                           GAACGTTCGAGATGA
                                                                                                                 22 BP; 6 A; 3 C; 7 G; 6 T; 0 other;
                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-188583P.
2001US-0802686.
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                                                                     71.4%; Score 15; DB 100.0%; Pred. No. 19;
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RESULT 24 AAS14665

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RESULT 25
AAH75992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Suppressing a respiratory syncytial virus infection by administering immunostimulatory sequence at the site of infection is useful to prevent and treat lower respiratory tract viral infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Respiratory syncytial virus. Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     respiratory syncytial virus; RSV; influenza virus; rhinovirus; adenovirus; measles virus; mumps virus; parainfluenza virus; rubella virus; poxvirus; parvovirus; hantavirus; varicella virus.
            Immunomodulatory oligonucleotide
                                         15-NOV-2001
                                                                  AAH75992;
                                                                                            AAH75992 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                       Sequence 22 BP; 6 A; 4 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 15; 40pp; English
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09-MAR-2001;
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                                       (first entry)
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2001US-0802686.
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            WO200168143-A2
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RESULT 26
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunomodulatory; immunostimulatory; Th1-type immune response; Th2-type immune response; interferon; idiopathic pulmonary fibrosis; viral infection; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                               Sequence
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09-MAR-2001;
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modified_base
Synthetic
                            Th2-type immune viral infection;
                                                                                                                      15-NOV-2001
                                                                                                                                                                               AAH75993 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunomodulatory polynucleotide/microcarrier complexes comprise an
immunostimulatory sequence containing polynucleotide linked to a
nonbiodegradable microcarrier -
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                                          Immunomodulatory; immunostimulatory; Th1-type immune response;
Th2-type immune response; interferon; idiopathic pulmonary fik
                                                                                       Immunomodulatory oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                            infection) and reducing levels of IgE.
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2001US-0802376.
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                                                                                                                                                                                                                                                                                                                                 71.4%;
                                                                                                                                                                                                                                                                                                                                                                                 3 C; 7 G; 6 T; 0 other;
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                                             fibrosis;
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RESULT 27
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Reducing risk of anaphylactic hypersensitivity response to in a subject, by administering an immunomodulating nucleic molecule comprising a specific sequence -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or suppressing a Th2-type response), increasing interferon-gamma (especially in a patient suffering from idiopathic pulmonary fibrosis), increasing interferon-alpha (especially in patients suffering from viral infection) and reducing levels of IgE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microcarrier complexes. The complexes comprise an immunostimulatory sequence (ISS), e.g. the present sequence, linked to a nonbiodegradable microcarrier provided that if the microcarrier is gold, latex or magnetic then the linkage is not biotin/avidin. The complex is useful for modulating an immune response (especially stimulating a Th1-type response
                                                     WPI; 2001-475812/51.
                                                                               Raz
                                                                                                                                21-DEC-1999;
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                                                                                                                                                                                                                                     Synthetic
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                                                                                                                                                                                                                                                                                                    Phosphorothioate
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                                                                                                        (REGC ) UNIV CALIFORNIA.
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15; Conserv
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2001US-0802376.
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                                                                                                                                                                                                                                                                                                   beta-gal/immunostimulatory oligonucleotide
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RESULT 28
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Best Local S
Matches 15
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                                       The present invention describes a method of treating an infection caused by an intracellular pathogen, involving administering to the patient an immunomodulatory nucleic acid and an anti-pathogenic agent. This is particularly useful in the treatment of mycobacterial infections. The present sequence is an immunomodulatory nucleic acid described in the
Sequence
                                                                                                                                                              Treating infection of intracellular pathogen e.g., Mycobacterium, in a subject, involves administering immunomodulatory nucleic acid molecule to inhibit intracellular replication of intracellular pathogen
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                                                                                                                                                                                                                                                                                                            31-JAN-2000; 2000US-0179353.
                                                                                                                                                                                                                                                                                                                                         30-JAN-2001; 2001WO-US03029.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          G3PDH gene; immunomodulatory oligonucleotide; infection; mycobacterium; intracellular pathogen; anti-pathogenic; ss.
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                             exemplification
                                                                                                                                 Examples; Page 26; 54pp; English.
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anaphylactic hypersensitivity or risk of anaphylactic response in
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A; 3 C; 7
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The present invention describes a composition (I) for intracellular delivery of a protein, comprising a protein in operative association with a cationic intracellular delivery vehicle comprising a cationic lipid, where the intracellular delivery vehicle is adapted to fuse with a cell membrane, therefore effecting intracellular delivery of the associated protein. Also described is a method for delivering a protein contact of the protein associated with a cationic lipid in such a manner so as to form an intracellular delivery composition, and contacting the delivery composition with a cell membrane and delivers the protein into the cell. (I) is useful in the preparation of a medicament for intracellular delivery of a therapeutic or prophylactic protein. (I) is useful for delivering antibodies to intracellular proteins to neutralise their activity, and contactuse. (I) is useful for medicament for intracellular cativity, and contactuse. (I) is useful for delivery of antibodies to intracellular proteins to neutralise their activity, and contactuse. (I) is useful for the in vitro or in vivo delivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific intracellular callivery of antibodies or peptides which block the function of specific in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Composition useful for intracellular delivery of a protein, comprises a protein in operative association with a cationic intracellular delivery vehicle comprising a cationic lipid, which is adapted to fuse with a
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The present invention describes immunomodulatory populations ((I) and (II) of conjugate molecules (CMs) comprising immunostimulatory sequences (ISS) of polymucleotides and antigens. The extent of conjugation affects the immunological properties (e.g. the extent of antigen-specific antibody formation, including Th1-associated antibody formation) so the conjugates are used for altering the type and extent of immune response. (I) and (II) have immunomodulatory, immunosuppressive and antiallergic activities, and can be used in the modulation of immune responses via the stimulation of Th1 lymphocytes and Th1-associated cytokines, and suppression of Th2 lymphocytes and Th1-associated cytokines, and (II) of conjugate molecules may be used for modulating immune responses in individuals e.g. for the treatment of an allergic condition. (I) and (II) may be used to modulate immune responses and therefore prevent potentially harmful reactions to antigens. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proteins and affect cellular metabolism, cell viability or virus replication. (I) is useful for delivering any protein of interest, including therapeutically useful proteins (e.g. tumour suppressor proteins, cystic fibrosis transmembrane regulator (CFTR), adenosine deaminase (ADA), hexoseaminidase A, peptides, wild type protein counterparts of mutant proteins and cell surface receptors) such as those for cytokines (e.g., interleuking, interferons, colony stimulating factors) and peptide hormones. The present sequence represents a peptide nucleic acid (PNA) oligonucleotide which is used in an example from the present invention for intracellular delivery of proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory sequence; ISS; immunomodulatory; immune response; antigen; antiallergic; modulation; Th1 lymphocyte stimulation; alle Th1-associated cytokine; Th2 lymphocyte suppression; cytokine; ss.
                                                                                                                                                                                                                                                                                                                                 controlling
                                                                                                                                                                                                                                                                                                                                             Populations of conjugate molecules comprising polynucleotide immunostimulatory sequences polynucleotides and antigens, useful for
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-329209/34.
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14-NOV-2000; 2000US-0713136
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Pred. No.
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RESULT 31
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                                                                                               (II) of conjugate molecules (CMs) comprising immunostimulatory sequences (CISS) of polynuclectides and antigens. The extent of conjugation affects the immunological properties (e.g. the extent of antigen-specific antibody formation, including Th1-associated antibody formation) so the conjugates are used for altering the type and extent of immune response. (I) and (II) have immunomodulatory, immunosuppressive and antiallergic activities, and can be used in the modulation of immune responses via the stimulation of Th1 lymphocytes and Th1-associated cytokines, and suppression of Th2 lymphocytes and cytokines. The populations (I) and (II) may be used to modulate may be used for modulating immune responses in individuals e.g. for the treatment of an altergic condition. (I) and (II) may be used to modulate immune responses and therefore prevent potentially harmful reactions to antigens. The present sequence represents an ISS polynuclectide which is used in the exemplification
Best Loca
Matches
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                                                          Sequence 22
                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 30; 97pp; English
                                                                                                                                                                                                                                                                                                                                                                        controlling
                                                                                                                                                                                                                                                                                                                                                                                    Populations of conjugate molecules comprising polynucleotide immunostimulatory sequences polynucleotides and antigens, us
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-329209/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tuck S,
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14-NOV-2000; 2000US-0713136.
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antigen; antiallergic; modulation; Th1 lymphocyte stimulation; alle
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               Local
                                                                                                                                                                                                                                                                                                               present invention describes immunomodulatory populations ((I) and
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l Similarity
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 Conservative
                                                          BP; 6 A; 4 C; 7 G; 5 T; 0 other;
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               71.4%; Score 15; 100.0%; Pred. No.
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RESULT 32
AAH20403
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                                             CC Unmethylated CG dinucleotide sequences are commonly found in bacterial CC DNA, and have been found to stimulate the innate immune system. Natural CK killer and T cells are activated by exposure to oligonucleotides CC containing CpG motifs. Oligonucleotides containing CpG motifs can be used CC as adjuvants in vaccines. The present invention relates to a CpG CC receptor. The CpG receptor contains a Toll homology domain (THD). The CC CpG receptor family are associated with responses to pathogens. CpG CC oligonucleotides may act as stimulators of various immune responses. The CC CpG receptor or cells expressing the receptor are useful for identifying CC compound which binds to or modulates an activity of the CpG receptor. CC CpH receptor dimmune responses antibacterials, (e.g. protection from CC ilsterial infection), tumour immunotherapy, allergy treatment, (e.g. suppressing LpB in human PBMC, shifting from Th2 to Th1) and as anti-inflammatory agents (e.g. for use in cystic fibrosis, sepsis, heart Cd isease, chlamydia, inflammatory bowel disease, arthritis and multiple sclerosis). The present sequence represents a CpG motif containing coligonucleotide used in examples demonstrating that CpG oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immune system stimulator; CpG motif; CpG receptor; CpG-R; antibacterial; immune response; vaccine adjuvant; tumour immunotherapy; allergy; anti-inflammatory; cystic fibrosis; sepsis; heart disease; chlamydia; inflammatory bowel disease; arthritis; multiple sclerosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                Novel CpG receptor and nucleic acid molecule encoding the receptor, for modulating immune response and for identifying compounds of therapeutic use which bind and/or modulate the activity of the receptor -
                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 14; 41pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mackichan ML;
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24-NOV-1999;
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                                 activate the MAPK pathways and NF-kappaB
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99US-0167389.
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Query Match

Sequence

22 BP;

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A; 3 C; 7 G; 6 T; 0 other;

Score 15;

DB 22;

Length 22;

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RESULT 34
AAF77041
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AC AAF77
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Best Local Similarity
Matches 15; Conserv
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     WO200112223-A2
                                                           Synthetic
                                                                                                           Modulate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a second antigen in an individual, involving administering to the individual an immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS) and a first antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 31; Page 15; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modulating immune response to a second antigen in humans involve administering an immunostimulatory polynucleotide comprising an immunostimulatory sequence and a first antigen -
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                                                                                                                                                                      Immunostimulatory DNA #1.
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                                                                                                             immune;
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AAF29800
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The present invention describes a method for enhancing an immune response to a substance, comprising administering an immunostimulatory nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulatory nucleotide sequence; antibody production; IFNgamma release; infection; allergy; ds.
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                                                                                 Example 1; Page 14; 47pp; English
                                                                                                                                 Enhancing immune response against pathogen or antigen associated with infectious diseases, an allergen or cancer, involves administering immunostimulatory nucleotide sequence prior to antigen exposure -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JUN-2000; 2000WO-US18229.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
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                                                                                                                                                                                                                                                                                                    Kobayashi H;
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CTL activity; Th1 response;
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                                                                                                                                         This invention describes a novel immunogenic composition (I), comprising a whole-killed human immunodeficiency virus (HIV) devoid of outer envelope protein gp120, an isolated nucleic acid molecule containing an immunostimulatory sequence (ISS) and an adjuvant, which enhances beta-chemokine levels in a mammal. The products of the invention have anti-HIV activity. (I) is useful for immunizing and for inhibiting AIDS in a mammal can be such as a human, (HIV seronegative or seropositive humans) or a rodent, in particular the primate is a pregnant mother or an infant. (I) can induce potent Th1 immune responses against a broad spectrum of HIV epitopes and provides a
                                                                                                     Sequence 22
                                                                                                                                                                                                                                                                                                       Example I; Page 26; 64pp; English.
                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus (HIV) compositions useful for immunizing and inhibiting AIDS in mammals, comprises HIV devoid of outer envelope protein and an immunostimulatory nucleic acid sequence .
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25-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunogenic; human immunodeficiency virus; immunostimulatory sequence; ISS; beta-chemokine; anti-HIV; AIDS; Th1 immune response; primer; HIV-specific cytotoxic T lymphocyte response; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence to a subject prior to exposure to the substance. This can be used to enhance antibody production, IRNgamma release, CTL activity at Thi related effects. The method can be used in the prevention and treatment of allergies, cancer and infections.
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GAACGTTCGAGATGA 20
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                                           Conservative
                                                                                                  BP; 6 A; 3 C; 7 G; 6 T; 0 other;
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Conservative 0;
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                                                                                                                                          The present invention describes an immunogenic composition (I) comprising a Neisseria antigen and an adjuvant composition comprising an oligonucleotide comprising at least 1 CG motif. Also described is an adjuvant composition (II) comprising an oligonucleotide which comprises at least 1 CG motif and a complete Freund's adjuvant (CFA), where the oligonucleotide preferably comprises at least one phosphorothicate bond. AAA92385 to AAA92385 represent specifically claimed oligonucleotides of the present invention. (I) is useful for stimulating an immune response in a mammal, preferably a human, against Neisseria infection, preferably Neisseria meningitidis infection and in the manufacture of a medicament
                                                                                                      Sequence
                                                                                                                                                                                                                                                                                          Claim 19; Page 9; 39pp; English.
                                                                                                                                                                                                                                                                                                                   Immunogenic composition useful for stimulating an immune response in a mammal against Neisseria infection, comprises Neisseria antigen and an adjuvant composition comprising an oligonucleotide with a CG motif -
                                                                                                                                                                                                                                                                                                                                                                                                     Grandi G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-FEB-2000; 2000WO-IB00176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neisseria antigen; Neisseria meningitidis; Neisseria gonorrhoeae; bactericidal; antibacterial; vaccine; immunostimulatory; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CG motif; complete Freund's adjuvant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CG motif and CFA containing oligonucleotide
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                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
                                                                                                      A; 3 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                               71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "preferably contains at phosphorothicate bond"
                                                                                                                                                                                                                                                                                                                                                                                                     Giuliani MM,
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                                                                                                                                immune response in a
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                                                               Score 15;
Pred. No.
                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                     Pizza
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                                                                 . 19;
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                                                                                                                                  mamma1.
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                                                                            Length 22;
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RESULT 38 ABV73190

ABV73190

standard; DNA;

22

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RESULT 39
ABQ78627
ID ABQ78
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AC ABQ78
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AC ABQ78
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AC ABQ78
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                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to suppressing symptoms of allergic response that involves administering to an antigen-sensitized mammalian host a dose of a composition comprising an immunomodulatory nucleic acid, and a second dose of a composition comprising an immunomodulatory nucleic acid, about 1 day - 8 weeks after the first dose. The immunomodulatory nucleic acid comprises a nucleotide sequence comprising 5'-CG-3'. The methods are useful for suppressing a symptom of an allergic reaction in a subject, maintaining suppression of a Th2 immune response and maintaining stimulation of a Th1 immune response. One method is useful in preventing the onset of, or rapidly suppress, antigen-stimulated inflammation associated with allergy, including antigenetimulatory puscased as occurs in the respiratory passages of asthmatics during an asthma attack, for boosting the immune respiratory passages of asthmatics during an asthma attack, for boosting the immune respiratory passages of asthmatics during an asthma attack.
                                                                                                                                                                                                                                                                                                                                                                                       the immune responsiveness of a mammalian host to a sensitizing antigen, and for treating a host suffering from inflammatory conditions such as asthma, nasal polyposis, allergic rinnitis, atopic dermatitis, allergic conjunctivitis, eosinophilic fasciltis, idiopathic hypereosinophilic conjunctivitis, eosinophilic fasciltis, idiopathic hypereosinophilic syndrome, and cutaneous basophil hypersensitivity. The present sequence represents the nucleotide sequence of an immunomodulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suppressing a symptom of an allergic response in a subject, useful for preventing inflammation associated with allergy, comprises administering to an antigen-sensitized host first and second doses of
                   25-NOV-2002
                                                      ABQ78627;
                                                                                         ABQ78627 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Examples; Page 27; 98pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-MAR-2001; 2001US-276865P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAR-2002; 2002WO-US08207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunomodulator; immunostimulant; antiinflammatory; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleotide sequence of an immunostimulatory oligonucleotide ISS-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunomodulatory nucleic acid
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                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                       GAACGTTCGAGATGA 20
                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                        GAACGTTCGAGATGA 22
                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 3 C; 7
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                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dermatological; vaccine; gene therapy; immune response; ss.
                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                71.4%;
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                                                                                                                                                                                                                                                                               Score 15; DB 24; 
; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                       G; 6 T; 0
                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                       other;
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RESULT 40
AAL44504
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents an exemplary immunostimulatory sequence (ISS) which enhances HIV-specific Thl cytckine and humoral responses, and also enhances both non-specific Thl cytckine and humoral responses, and also enhances both non-specific and HIV-specific beta-chemokine production. ISSs can be included in HIV immunogenic compositions of the invention. The specification describes a method for treating an HIV-infected individual, which comprises combining immunisation with an anti-retroviral treatment and withdrawal from treatment. The cycles of anti-retroviral treatment and withdrawal from treatment. The advantages of the method of the invention include a delay in the rebound to an unacceptably high viral load; a more rapid or sustained increase in HIV-specific CD4 T cell counts; a reduction or delay in the development of AIDS symptoms, including AIDS-related opportunistic infections; and a higher degree of patient compliance with treatment and fewer toxic side effects associated with long-term anti-retroviral directions treatment. The
                                                               Vaccine; immune response; microparticle; ds; adsorbent surface; poly(alpha-hydroxy acid); polyhydroxy butyric acid; polycaprolactone; polyorthoester; polycyanacrylate; detergent; submicron emulsion; viral infection; bacterial infection; parasitic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating an HIV-infected individual comprises treatment with anti-retroviral compound and immunization with an HIV immunogenic composition with structured cycles of anti-retroviral treatment and
                                                                                                                                                         CpG motif oligonucleotide #12.
                                                                                                                                                                                            08-NOV-2002
                                                                                                                                                                                                                                                             AAL44504 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     withdrawal from treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-2002; 2002WO-US02077.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulatory sequence; ISS; Th1 cytokine response; humoral response;
HIV; beta-chemokine; immunisation; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                        15;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                      GAACGITCGAGAIGA 20
                                                                                                                                                                                                                                                                                                                                                       GAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 BP; 6 A; 3 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 15; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for treating an HIV-infected
                                                                                                                                                                                                                                                                                                                                                                                                                                         71.4%; Score 15; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                     22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Th1 cytokine and humoral responses.
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Unidentified

CpG oligonucleotide

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RESULT 41
ABQ75153
ID ABQ75
XX ABQ75
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of raising an immune response in a host animal. The method of the invention comprises administering a microparticle that has an adsorbent surface to which a first biologically active macromolecule (e.g. a nucleic acid) has been adsorbed. The microparticle comprises a polymer microparticle of poly(alpha-hydroxy acid), a polyhydroxy butyric acid, a polycaprolactone, a polyorthoester, a polycaprolactone, a polyorthoester, a polycaprolactone of the invention is useful for immunising a host animal against viral, bacterial or parasitic infections. The present DNA
                                                                                                                                                                                   misc_RNA
                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                          Immunostimulatory sequence; ISS: immunomodulatory; immune response; allergy; asthma; infectious disease; interferon-gamma; IFN-gamma; idiopathic pulmonary fibrosis; viral infection; mycobacterial disease; malaria; leishmaniasis; toxoplasmosis; schietosomiasis; clonorchiasis; immunoglobulin B; IgB-related disorder; antiallergic; antiasthmatic; virucide; antibacterial; protozoacide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A microparticle to which a biologically active macromolecule is adsorbed, for use as a vaccine composition to treat viral, bacterial parasitic infections, comprises a polymer microparticle, a detergent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-519084/55
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30-AUG-2001;
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                           04-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ISS immunomodulatory oligonucleotide SEQ ID NO:2.
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T. Dubensky
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15; Conserv
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2001US-315905P
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/note= "uracil"
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Pred. No.
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19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bacterial or
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RESULT 42
ABQ75163
ID ABQ75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC comprising an immunostimulatory sequence (ISS). Also described: (I) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and comprising (I). (I) has antiallergic, antiasthmatic, virucide, (3) a kit comprising (I); (I) has antiallergic, antiasthmatic, virucide, (2) a minume response (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune componse, especially an allergy or asthma, or an infectious disease. (I) is also useful for increasing interferon-gamma (IFN-gamma) in an comprising a symptom of an infectious disease, or IFN-alpha in an complicating a symptom of an infectious disease caused by a cellular cathogen such as mycobacterial disease, malaria, leishmaniasis, complements and complements in an individual, or a symptom of an immunoglobulin E (IgD)-related disorder, preferably an complements and immunomodulatory oligonucleotide which compresent sequence represents an immunomodulatory oligonucleotide which complements and the present invention.
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunomodulatory polynucleotide for modulating an immune response in subject suffering from disorders associated with Th2-type immune response, e.g. allergy, or infectious disease, comprises an immunostimulatory sequence
                                                                                                             Synthetic.
                                                                                                                                                                                                 Immunostimulatory sequence; ISS: immunomodulatory; immune response; allergy; asthma; infectious disease; interferon-gamma; IFN-gamma; idiopathic pulmonary fibrosis; viral infection; mycobacterial disease;
                                                                                                                                                                                                                                                                                                                                                                                          ABQ75163 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes an immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS). Also described: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 20; 95pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-DEC-2000; 2000US-258675P.
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                                                                        WO200252002-A2
                                                                                                                                               virucide; antibacterial; protozoacide; ss.
                                                                                                                                                                                   malaria; leishmaniasis; toxopl
                                                                                                                                                                                                                                                                             ISS immunomodulatory oligonucleotide SEQ ID NO:12.
                                                                                                                                                                                                                                                                                                                  05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                    ABQ75163;
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                                                                                                                                                                  mmunoglobulin E; IgE-related disorder; antiallergic;
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Pred. No. 19;
                                                                                                                                                                                   asmosis; schistosomiasis; clonorchiasis;
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27-DEC-2001; 2001WO-US50821.

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    individual suffering from disorders associated with a Th2-type immune response, especially an allergy or asthma, or an infectious disease. is also useful for increasing interferon-gamma (IFM-gamma) in an individual having idiopathic pulmonary fibrosis, or IFM-alpha in an individual having a viral infection. (I) is further useful for ameliorating a symptom of an infectious disease caused by a cellular pathogen such as mycobacterial disease, malaria, leishmaniasis, toxoplasmosis, schistosomiasis and clonorchiasis in an individual, or symptom of an immunoglobulin E (IgE)-related disorder, preferably an allergy-related disorder, in particular asthma in an individual. The present sequence represents an immunomodulatory oligonucleotide from
                                                                                                                                                                               Immunostimulatory sequence; ISS: immunomodulatory; immune response; allergy; asthma; infectious disease; interferon-gamma; IFN-gamma; idiopathic pulmonary fibrosis; viral infection; mycobacterial disease; malaria; leishmaniasis; toxoplasmosis; schistosomiasis; clonorchiasis; immunoglobulin E; IgE-related disorder; antiallergic; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising an immunostimulatory sequence (ISS). Also described: (1) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I); (2) an immunomodulatory polynucleotide/microcarrier (IMPMC) complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and (3) a kit comprising (I). (I) has antiallergic, antiasthmatic, virucide, antibacterial and protozoacide activities, and can be used as a modulator of immune response. (I) is useful for modulating an immune response in an individual suffering from disorders associated with a Th2-type immune (1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
               27-DEC-2001; 2001WO-US50821.
                                                       04-JUL-2002.
                                                                                            WO200252002-A2
                                                                                                                               Synthetic
                                                                                                                                                                     virucide;
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                                                                                                                                                                                                                                                                                                                                                                                                                 ABQ75173 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 5 A; 4 C; 7
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                                                                                                                                                                                                                                                                                                   ISS immunomodulatory oligonucleotide SEQ ID NO:22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention describes an immunomodulatory polynucleotide (I) prising an immunostimulatory sequence (ISS). Also described: (1) an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-657426/70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAACGTTCGAGATGA 22
                                                                                                                                                                   antibacterial; protozoacide;
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100.0%; Pr
                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC comprising an immunostimulatory sequence (ISS). Also described: (I) an CC immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition comprising (I); (2) an immunomodulatory composition complex, complex, comprising (I) linked to a biodegradable MC, where the MC is less than 10 micrometre in size; and CC distance in an immunomodulatory composition (I) and immunomodulatory composition (I) is useful for modulating an immune response in an CC individual suffering from disorders associated with a Th2-type immune cresponse, especially an allergy or asthma, or an infectious disease. (I) is also useful for increasing interferon-gamma (IFN-alpha in an CC individual having a dispathic pulmonary fibrosis, or IFN-alpha in an CC individual having a viral infection. (I) is further useful for CC ameliorating a symptom of an infectious disease caused by a cellular CC pathogen such as mycobacterial disease, malaria, leishmaniasis, and CC complexmosis, schistosomiasis and clonorchiasis in an individual, or a CC symptom of an immunoglobulin E (IgE) related disorder, preferably an CC present sequence represents an immunomodulatory oligonucleotide from the present sequence represents an immunomodulatory oligonucleotide from the present sequence represents an immunomodulatory oligonucleotide from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                    allergy; asthma; idiopathic pulmor
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                                                                                                                                                           Synthetic
                                                                                                                                                                                                    virucide;
                                                                                                                                                                                                                                          malaria;
                                                                                                                                                                                                                                                                                                                                         ISS immunomodulatory oligonucleotide SEQ ID
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                                                                                                 misc_RNA
                                                                                                                                                                                                                       immunoglobulin
                                                                                                                                                                                                                                                                               Immunostimulatory sequence; ISS: immunomodulatory; immune response;
allergy; asthma; infectious disease; interferon-gamma; IFN-gamma;
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                                                                                                                                                                                          asthma; infectious disease; interferon-gamma; IFN-gamma; oc pulmonary fibrosis; viral infection; mycobacterial disease; leishmaniasis; toxoplasmosis; schistosomiasis; clonorchiasis; bulin E; IgE-related disorder; antiallergic; antiasthmatic; antibacterial; protozoacide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
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                                                                                                     Location/Qualifiers
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RESULT 45
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                                                                                                                                                                                                         Immunostimulatory sequence; ISS: immunomodulatory; immune response; allergy; asthma; infectious disease; interferon-gamma; IFN-gamma; idiopathic pulmonary fibrosis; viral infection: mvcohacterial Aiaco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunomodulatory polynucleotide for modulating an immune response subject suffering from disorders associated with Th2-type immune response, e.g. allergy, or infectious disease, comprises an immunostimulatory sequence
                                                                                             Synthetic.
                                                                                                                                        virucide;
                                                                                                                                                                                                                                                                                                         ISS immunomodulatory oligonucleotide SEQ ID NO:45
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                       modified_base
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                                                                                                                                                              leishmaniasis; toxoplasmosis; schistosomiasis; obulin E; IgE-related disorder; antiallergic; ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAACGITCGAGAIGA 20
                                                                                                                                                                                      sthma; infectious disease; interferon-gamma; IFN-gamma; pulmonary fibrosis; viral infection; mycobacterial disease; eishmaniasis; toxoplasmosis; schistosomiasis; clonorchiasis;
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                                                                                                                                        protozoacide;
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Best Local
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Immunostimulatory sequence; ISS: immunomodulatory; immune response; allergy; asthma; infectious disease; interferon-gamma; IFN-gamma; idiopathic pulmonary fibrosis; viral infection; mycobacterial disea
                                                                                                              05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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15; Conserv
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'note= "5-bromocytosine"
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== "5-bromocytosine"
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RESULT 47
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Best Local :
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             05-NOV-2002
                                              ABQ75259;
                                                                              ABQ75259 standard; DNA; 22
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            (first entry)
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RESULT 48
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                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence represents an immunomodulatory related oligonucleotide which was used in an example from the present invention.
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                                                                                                                         AAD24885
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 6 A; 3 C; 7 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                             Conservative
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12-MAR-2002

(first entry

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Immunostimulatory oligodeoxynucleotide (ISS-ODN)

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CC The invention relates to a method for modulating cell death or reducing CC DNA damage in an eukaryotic cell by contacting the cell with an agent CC (DNA-PK). The invention also relates nucleic acids which modulate the CC (DNA-PK). The invention also relates nucleic acids which modulate the CC immune response binding to Ku antigen, resulting in activation of DNA-PK. CC The method is useful for modulating cell death or reducing DNA damage in CC an eukaryotic cell, for treating any disorder resulting from a genotoxic CC insert to a cell e.g., necrosis, apoptosis. The method is also useful CC retreating cell death-related indications such as Alzheimer's disease, CC central nervous system inflammation, osteoporosis, degenerative liver CC disease, crebbellar degeneration, reperfusion injury, multiple sclerosis, cc amyotrophic lateral sclerosis, myocardial infarction, head injury damage, acquired immunodeficiency syndrome (AIDS), aplastic anaemia, cerebral CC infarction, bypass heart surgery, organ transplantation. The method is also useful for treating follicular lymphomas, carcinomas, autoimmune CC alsorders (systemic lupus erythematosus), hormone dependent tumours, cused for identifying ISS-binding protein, which is used in the CC exemplification of the invention.
                                                     RESULT 49
AAS16337
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                                    AAS16337 standard; DNA;
                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modulating cell death or reducing DNA damage in eukaryotic cells, useful for reducing cell death in individual or organ, comprises contacting cell with agent modulating biological activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
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17-JAN-2001;
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2001US-262321P.
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100.0%;
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AAS16338 standard; DNA; 22

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RESULT 50
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Note: The present sequence is shown as single stranded in the specification, but the patentees state on page 20 that this sequence may
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Preventing, reducing the severity or reducing the recurrence of an infection or symptom of herpes simplex virus (HSV), e.g. HSV-2, comprises administering an immunostimulatory sequence to an individual
                                                                                                                                                                                                                                                                                                                                                                            Sequence
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09-MAR-2001;
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15; Conserv
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RESULT 51
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       labialis (Cold sores) and genital herpes. The present sequence one of the ISS polynucleotides of the invention.

Note: The present sequence is shown as single stranded in the specification, but the patentees state on page 20 that this se
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Animal papillomavirus infection; human papillomavirus; HPV; STD; wart; sexually transmitted disease; cervical cancer; immune response;
                                                                                                     ISS polynucleotide #1 useful for treating papillomavirus infections.
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09-MAR-2001;
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2001US-0802518.
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100.0%; Prr
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RESULT 52
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Best Local (
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09-MAR-2001;
                        Animal papillomavirus infection; human papillomavirus; HPV; STD; wart; sexually transmitted disease; cervical cancer; immune response;
                                                                                                    14-FEB-2002
                                                                                                                                AAS16349;
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                                                                                                                                                                                                                                                                                                                                                                                       be double stranded
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             immunostimulatory
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                                                                    polynucleotide
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                                                                                                                                                            standard; DNA; 22
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2001US-0802445
                                                                                                 (first entry)
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          sequence;
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                                                                     #2 useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         omposition comprising a polynucleotide sequence to the individual -
             iss;
                                                                    for treating
             virucide;
                                                                                                                                                                                                                                                                                                              DB 24;
. 19;
                                                                    papillomavirus infections.
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RESULT 53
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Best Local S
Matches 15
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                                                                         Cytotoxic T lymphocyte; CTL; T cell; tumour load; cancer radiotherapy; immunostimulatory sequence oligonucleotide; ISS-ODN; chemotherapy; immunosuppression; transplantation; autoimmune disease; infection; acquired immune deficiency syndrome; AIDS; intracellular pathogen; cytomegalovirus; mycobacterial infection; Epstein-Barr virus; varicella zoster virus; human immunodeficiency virus; HIV; phosphorothioate backbone; ss.
modified_base
                                               Unidentified
                                                                                                                                                                                                  Immunostimulatory sequence oligonucleotide (ISS-ODN) #1.
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09-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double stranded
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                                                                                                                                                                                                                                                                                                                                                                       GAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                  GAACGITCGAGATGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  preventing or ameliorating papillomavirus infections, administering a composition comprising a polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunostimulatory sequence to
                                                                                                                                                                                                                                                                                                                                                                                                                               71.4%; Score 15; DB 24; llarity 100.0%; Pred. No. 19; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A;
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2001US-0802445.
                                                                                                                                                                                                                                (first entry)
 Location/Qualifiers
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RESULT 54
ABA03833
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                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method for increasing antigen-specific CC cytotoxic T lymphocyte (CTL) activity in a CD4+ T cell-deficient CC individual, comprising administering an immunostimulatory sequence CC oligonucleotide (ISS-ODN). The immunostimulatory nucleic acids of the CC invention are used in CD4+ T cell-deficient individuals to decrease CC tumour load, to treat a primary or acquired immunodeficiency; the control of the CC particularly where the acquired immunodeficiency is temporary and due CC concer radiotherapy or chemotherapy or immunosuppression following CC bone marrow or organ transplantation, or autoimmune disease treatment, CC or is acquired immunodeficiency syndrome (AIDS). The nucleic acids may CC used to treat a person at risk of becoming CD4+ T cell-deficient, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC used to treat infection, particularly by an intracellular pathogen, CC disulfide-linked phosphorothioate immunostimulatory sequence is a 5' CC coligonucleotide (ISS-ODN), used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                  Matches
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Best Local &
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11-MAY-2000;
05-JUL-2000;
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(VETE-)
                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Increasing antigen-specific cytotoxic T lymphocyte activity in a CD4+cell deficient individual, useful to treat immunodeficiency and block
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-010699/01.
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Immunomodulatory polynucleotide/microcarrier complex; IMP/MC;
                              Immunostimulatory sequence (ISS)
                                                                                                  ABA03833;
                                                                                                                               ABA03833 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV infection, comprises administering immunostimulatory nucleic acid
                                                                12-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                 22 BP;
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; 2000US-203567P.
; 2000US-215895P.
                                                                                                                                                                                                                                                                                  Conservative
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VETERANS AFFAIRS
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                                                                (first entry)
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                 Score 15;
Pred. No.
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                                  SEQ ID NO:1.
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19;
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microcarrier (MP/MC) complex (I), which comprises a polynucleotide comicrocarrier (MC). The ISS comprises the sequence: 5'-G-3', where the comicrocarrier (MC). The ISS comprises the sequence: 5'-G-3', where the comicrocarrier (MC). The ISS comprises the sequence: 5'-G-3', where the CC antiallergic, antibacterial, antiprotozoal, antiparasitic, hepatotropic and nephrotropic activities. It can be used as an interferon (IFM)-alpha stimulator, IFN-gamma stimulator or an immunoglobulin E (IgE) stimulator. (I) can be used for modulating the immune response of individuals, cutaneous creatisticn-induced fibrosis (IPF), scleroderma, cutaneous creatisticn-induced fibrosis, hepatic fibrosis, including schistosomiasis-induced hepatic fibrosis, renal fibrosis, infectious diseases caused by cellular pathogen (e.g. a mycobacterial disease, complex is beighned by cellular pathogen (e.g. a mycobacterial disease, complex is complex is complex is a microbacterial disease, complex is a microbacterial disease, complex is a mycobacterial disease, complex is mindividuals suffering from cancer, or in individuals at the mycobacterial disease is a specifically claimed ISS which can be used in an IMP/MC complex of the mycobacterial individuals suffering from cancer, or in individuals
                                                                                                                                                                Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New immunomodulatory polynucleotide/microcarrier complex, useful for modulating the immune response of individuals, particularly humans, for treating idiopathic pulmonary fibrosis, scleroderma, malaria or
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09-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 49; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         invention.
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GAACGTTCGAGATGA 22
                                                                       GAACGTTCGAGATGA 20
                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                         BP; 6
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2001US-0802359.
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                                                                                                                                                                                              71.4%;
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6 GAACGTTCGAGATGA 20

Query Match Best Local S Matches 15

Similarity

71.4%;

Score 15; Pred. No. Mismatches

DB 19;

24; 0

Length 22; Indels

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Gaps

0

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RESULT 55
                                                          CC microcarrier (IMP/MC) complex (I), which comprises a polynucleotide (C) having an immunostimulatory sequence (ISS) linked to a biodegradable microcarrier (MC). The ISS comprises the sequence: 5'-CG-3', where the CC microcarrier (MC). The ISS comprises the sequence: 5'-CG-3', where the CC microcarrier (MC). The ISS comprises the sequence: 5'-CG-3', where the CC microcarrier (MC) is less than 10 micro m in size. (I) has immunomodulator, and interferon (IFW)-alpha critical method of the complex is an interferon (IFW)-alpha cc stimulator. IFW-gamma stimulator or an immunoglobulin E (IgE) stimulator. (I) can be used as an interferon (IFW)-alpha cc particularly humans. The IMP/MC complex is particularly useful for treating idiopathic pulmonary fibrosis (IPP), scleroderma, cutaneous cradiation-induced fibrosis, hepatic fibrosis including radiation-induced fibrosis, hepatic fibrosis infectious consists complex is schistosomiasis or radiation-induced depatic fibrosis, renal fibrosis, infectious consists caused by cellular pathogen (e.g. a mycobacterial disease, caused by cellular pathogen (e.g. a mycobacterial disease, consortial, leighmaniasis, toxoplasmosis, schistosomiasis or chlaria, leighmaniasis, toxoplasmosis, schistosomiasis or chlaria, leighmaniasis, toxoplasmosis, schistosomiasis or prophylactic cresponse (e.g. aller be used in individuals receiving therapeutic or prophylactic critic of exposure to an infectious agent. The present sequence represents and the present sequence represents consorted to the present sequence represents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunomodulation; immunostimulation; phosphorothioate; immunomodulator; antiallergic; antibacterial; antiprotozoal; antiparasitic; hepatotropic; nephrotropic; interferon-alpha stimulator; interferon-gamma stimulator; immunoglobulin B stimulator; immune response; IPF; scleroderma; malaria;
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New immunomodulatory polynucleotide/microcarrier complex, useful for modulating the immune response of individuals, particularly humans, of creating idiopathic pulmonary fibrosis, scleroderma, malaria or
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09-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mycobacterial disease; toxoplasmosis; schistosomiasis; chlonorchiasis; allergy; allergy-induced asthma; prophylactic vaccine; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         idiopathic pulmonary fibrosis; cutaneous radiation-induced fibrosis; hepatic fibrosis; renal fibrosis; infectious disease; leishmaniasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 18; 63pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunostimulatory sequence (ISS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mmunomodulatory polynucleotide/microcarrier complex; IMP/MC; IgE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention describes an immunomodulatory polynucleotide/
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    22 BP; 6
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2001US-0802359.
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                                               the exemplification of the present
  A; 4 C; 7 G; 5 T; 0 other;
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                                            invention.
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RESULT 56
ABA03844
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Matches
                                                                         Query Match
                                                                                                                        The present invention describes a method for reducing severity of a symptom of virus infection in an individual infected with a virus. The method comprises administering a composition consisting of a polynucleotide having an immunostimulatory sequence (ISS). The ISS comprises the sequence 5'-C',0',pyrinidine,pyrinidine,C',G-3'. An antigen is administered in conjunction with the composition. ISS has virucide activity and can be used in gene therapy. The method using the ISS can be used for suppressing, ameliorating and/or preventing viral infections to an individual who may be at risk of being exposed to, exposed to or infected by a virus. It may also be used in reducing the recurrence or duration of a symptom of viral infection, delaying the development of a virus infection, and reducing viraemia or blood levels of virus antigens. The present sequence represents a specifically claimed ISS for use in the method of the invention.
                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                          Claim 4;
                                                                                                                                                                                                                                                                                                                                                   sequence
                                                                                                                                                                                                                                                                                                                                                             Reducing severity, recurrence or duration of symptom of virus infection, or reducing viremia or blood levels of virus antigen, comprises administering a polynucleotide having an immunostimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-048999/06
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09-MAR-2001;
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GAACGTTCGAGATGA 22
                        GAACGTTCGAGATGA 20
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                                                                                                    BP; 6
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2001US-0802685.
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/mod_base= OTHER
                                                                                                    A.
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                                                                                                    3 C;
                                                                           71.4%;
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                                                 score 15; DB; Pred. No. 19; 0; Mismatches
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                                                                                                    G; 6 T; 0 other;
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therapy; viraemia; phosphorothioate;
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                                                             DB 24;
19;
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RESULT 58
ABA03856
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ABA03845
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-MAR-2000;
09-MAR-2001;
                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                       exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method for reducing severity of a symptom of virus infection in an individual infected with a virus. T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprises administering a polynucleotide having an immunostimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reducing severity, recurrence or duration of symptom of virus infection, or reducing viremia or blood levels of virus antigen,
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         Immunostimulatory sequence (ISS) SEQ ID NO:1
                                    12-FEB-2002
                                                                                       ABA03856 standard;
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immunomodulation; virucide;
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                                                                                                                                                        GAACGTTCGAGATGA
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2001US-0802685.
                                   (first entry)
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                                                                                       DNA; 22
                                                                                                                                                                                                       71.4%; 50,
100.0%; Pr
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                                                                                                                                                                                                                                                                                         invention.
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therapy; viraemia;
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19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            levels of a hepatitis virus antigen in an individual infected with hepatitis B virus (HBV). The method comprises administering a composition comprising a polynucleotide having an immunostimulatory sequence (ISS) to the individual, where the ISS comprises the sequence 5'-C, G-3', an HBV antigen is not administered in conjunction with administration of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reducing viremia and blood levels of hepatitis virus antigen in an individual infected with hepatitis B virus, comprises administering composition comprising a polynucleotide having an immunostimulatory
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09-MAR-2001;
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                                                                                                                                         Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 5; Page 38; 43pp; English
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                                                                                     Local
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                                                                                     Similarity
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                  GAACGTTCGAGATGA 20
GAACGTTCGAGATGA
                                                                                                                                         BP; 6
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2001US-0802370.
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                                                                                     Score 15; DB 24; Pred. No. 19;
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RESULT 59
ABA03857
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ABA03857 standard; DNA;

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RESULT 60
AAS15592
ID AAS15
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AC AAS15

AAS15592 standard; DNA; 22

AAS15592;

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the composition, and where the composition is administered in an amount sufficient to reduce HBV viraemia or blood levels of a hepatitis virus antigen. ISS has virucide, anti-inflammatory and hepatotropic activities, and can be used in gene therapy. The method can be used for suppressing and/or ameliorating hepatitis infection in an individual, especially for preventing, palliating, ameliorating, reducing and/or eliminating one or more symptoms of HBV or HCV (hepatitis C virus) infection without administering HBV or HCV antigens. The method is specifically useful for reducing viraemia and hepatitis viral antigen in blood. ISS-containing polynucleotides may also be used to improve physical symptoms such as jaundice, fatigue, abdominal pain, and other clinical/ laboratory portal hypertension, or cirrhosis. The present sequence represents an invertion of the present of the present and the present sequence represents and invertion.
                                                                                                                   Matches
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                                                                                                                                                                                                                                      Sequence 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reducing viremia and blood levels of hepatitis virus antigen in an individual infected with hepatitis B virus, comprises administering composition comprising a polynucleotide having an immunostimulatory
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09-MAR-2001;
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GAACGITCGAGATGA 22
                                                        GAACGTTCGAGATGA 20
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2001US-0802370.
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                                                                                                                   Score 15; DB 24; ; Pred. No. 19; 0; Mismatches (
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                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a polynucleotide vaccine (I) comprising a concleic acid sequence encoding an antigen derived from a non-host species of a first phylum or first kingdom, where the nucleic acid sequence encoding the antigen is modified by deletion of a native signal sequence, and/or an immunomodulatory nucleic acid sequence. (I) is useful for modulating an immune response to an antigen, especially a plant (ragweed or grass pollen), food, latex, cat dander, cockroach or house dust mite allergen. (I) is also useful for eliciting an immune response to an antigen derived from a pathogen, such as bacterium, virus or a parasite. The vaccine is co-administered with an immunostimulatory nucleotide sequence which comprises an unmethylated 5'-CG-3' nucleotide sequence. Antigens of pathogens can be present sequence represents fungal, yeast or other pathogens. The present sequence represents with (I) to amplify the immune response to the co-administered allergen.
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                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Immunomodulatory composition; immune response; Th2-type immune response; allergy; asthma; infectious disease; vaccine; cancer; IFN-alpha; IFN-gamma; idiopathic pulmonary fibrosis; scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JAN-2002
                                                                       Nucleotide sequence of a control immunostimulatory oligonucleotide
                                                                                                         03-JUN-2003
                                                                                                                                                                              ABZ77582 standard;
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotide vaccine for eliciting immune response to an antigen derived from a pathogen, plant or food, comprises antigen-encoding nucleic acid sequence derived from non-host species of first phylum or
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                                                                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                   6
                                                                                                                                                                                                                                                                                                                                      l Similarity
15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                           22
                                                                                                                                                                                                                                                                                                     GAACGITCGAGATGA 20
                                                                                                                                                                                                                                                                                                                                                                                                         BP; 6 A;
                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                         (first entry)
                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                           3 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                        71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nguyen
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                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                        Score 15; DB 24; 
; Pred. No. 19;
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RESULT 62
AAL51531
ID AAL55
XX
AC AAL51
XX
DT 10-AI
XX
DE CTL 1
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                                                                                                                                                                                                                                                                                                                                                                                                      CC comprises a complex of a microcarrier and an immunocollatory

CC comprises a complex of a microcarrier and an immunocollatory

CC comprises a complex of a microcarrier and an immunocollatory

CC comprises a complex of a microcarrier and an immunocollatory

CC observed as the composition is useful for modulating an immunocollatory

CC response in an individual suffering from a disorder associated with a

CC rh2-type immune response which is selected from allergies and

CC rh2-type immune response which is selected from allergies and

CC receives a prophylactic or therapeutic vaccine, where the therapeutic

CC vaccine is selected from an epitope which is selected from an allergy

CC epitope, a mycobacterial epitope, which is selected from an allergy

CC epitope, a mycobacterial epitope, which is selected from an allergy

CC individual is at a risk of exposure to an infectious agent or at risk

CC fdeveloping cancer, or suffers from cancer or an infectious disease.

CC The composition is useful for increasing interferon (IFN)-gamma or

CC relicated from idiopathic pulmonary fibrosis, and renal fibrosis, viral

CC reditation-induced fibrosis, hepatic fibrosis, and renal fibrosis, viral

CC reditation-induced fibrosis caused by a cellular pathogen. The infectious

CC the infectious disease is caused by a cellular pathogen. The infectious

CC tooplasmosis, schistosomiasis, and clonorchiasis. The present sequence

CC represents an immunostimulatory oligonucleotide, used as a control in
                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunomodulatory composition for modulating immune response and reducing levels of immunoglobulin E in an individual having the disorder, has a complex of microcarrier and an immunomodulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cutaneous radiation-induced fibrosis; hepatic fibrosis; renal fibrosis; viral infection; mycobacterial disease; malaria; leishmaniasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-2001; 2001US-310743P.
25-OCT-2001; 2001US-335263P.
                                      10-APR-2003
                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The specification describes an immunomodulatory composition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 62; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-248018/25.
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                                                                           AAL51531;
                                                                                                              AAL51531 standard; DNA; 22 BP
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                                                                                                                                                                                                                                                                                l Similarity
15; Conserv
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                                                                                                                                                                                                        GAACGTTCGAGATGA 22
                                                                                                                                                                                                                                                                                                                                                                                              of the invention.
                                                                                                                                                                                                                                                                                                                                                       BP;
                                                                                                                                                                                                                                                                                Conservative (
                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                         A; 3 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                              71.4%; Score 15; DB
100.0%; Pred. No. 19;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                   DB 25;
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                                                                                                                                                                                                                                                                                                                     Length 22;
                                                                                                                                                                                                                                                                                    Indels
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CTL recognition antigen-related oligonucleotide,

SEQ ID

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RESULT 63
ABZ57964
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises a method for screening a cytotoxic T-lymphocyte (CTL) recognition antigen, which includes CTLs with antitumour effect against human T-lymphotrophic leukaemia virus-1 (HTLV-1) tumours. The CTL-recognition antigens identified by the method of the invention are useful as immune response-inducible vaccines, and as components of drug preparations and health foods. The present DNA sequence represents an objection of the inventional contents of the present of the contents of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Screening of cytotoxic T-lymphocyte-recognition antigen with a human T-lymphotrophic leukemia virus-1 (HTLV-1) antitumor effect, for use a vaccine, comprises administering a test substance to a
                                                                                                                                                                                                             modified_base
25-JUN-2002; 2002WO-IL00507.
                                                 03-JAN-2003
                                                                                              WO2003000232-A2
                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory oligodeoxynucleotide ISS-ODN 1018.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ57964;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ57964 standard; DNA; 22 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide that was used in an example of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HTLV-1-associated disease animal model
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                                                                                                                                                                                                                                                                                                                                                             ISS-ODN 1018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                 therapy;
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                                                                                                                                                                                                                                                                                                                                                   immunostimulant; vaccine; adjuvant; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                 liposome; ss
                                                                                                                                                                                                                Location/Qualifiers
1..22
                                                                                                                                                                                            *tag=
                                                                                                                                             note=
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100.0%; Pr
                                                                                                                                                                   base= "OTHER"
                                                                                                                                          "phosphorothioate linkage"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15;
Pred. No.
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19;
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RESULT 64
AAV80098/c
ID AAV800
XX AAV800
XX INMUNO
XX IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothicate or phosphodiester backbone. Liposomal ISS-ODN can be used e.g. as a vaccine adjuvant against pathogens and cancer, in the treatment or prevention of diseases caused by certain infectious microorganisms, in the treatment or prevention of allergic diseases, or to boost innate immunity. In examples of the invention, ISS-ODN 1018 was encapsulated in large multilamellar liposomes with up to 95% efficiency. The liposomal formulation was a considerably more potent parenteral adjuvant in mice than the soluble form of ISS-ODN, as shown in experiments with an influenza vaccine. Enhancement of the systemic humoral and cellular response was demonstrated by liposomal ISS-ODN 1018 co-administered with tuberculosis vaccine. Liposomal ISS-ODN 1018 was administered with tuberculosis vaccine. Liposomal ISS-ODN 1018 was also used as a adjuvant for a cancer (murine mammary carcinoma) vaccine, and activated resistance to Leishmaniasis when administered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
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                                                                Synthetic.
                                                                                                                      ISS: cancer; allergy; asthma; hepatitis B infection; papillomavirus; human immunodeficiency virus; influenza; herpes; M. tuberculosis; ss; B. pertussis; malaria; plasmodia; Leishmania; Trypanosoma; Schistosom
                                                                                                                                                                                                                                                                              Immunomodulatory oligo comprising an ISS sequence.
                                                                                                                                                                                                                                                                                                                                                                                                              AAV80098;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV80098 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 19; 68pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barenholz Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-JUN-2001; 2001US-300072P.
17-DEC-2001; 2001US-339785P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             σ
                                                                                                                                                                                                                       immunostimulatory;
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Mismatches 0
                                                                                                                                                                                                                       octanucleotide; immune regulation;
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AAA38067/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to immunomodulatory oligonucleotides that comprise at least 1 immunostimulatory octanucleotide sequence (ISS) where the ISS sequences are selected from the group consisting of AACGTTCC, AACGTTCG, GACGTTCC, and GACGTTCG. The immunomodulatory sequences are used to treat patients needing immune regulation, such as those suffering from cancer, an allergic disease and asthma. They are also used to prevent infectious disease such as influenza, herpeatitis B, human immunodeficiency and papillomavirus, Hemophilus influenza, Mycobacterium tuberculosis and Bordetella pertussis, malarial plasmodia, Leishmania, Trypanosoma and Schistosoma. The immunomodulatory sequences are used to screen for human schistosoma. The immunomodulatory sequences are used to screen for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schistosoma. The immunomodulatory sequences are used to screen for hum. immunostimulatory activity by incubating macrophage cells and the oligonucleotide; and determining the relative amount of Thi-biased cytokines in the supernatant. Sequences AAV80096 to AAV80103 represent specific claimed examples of such immunomodulatory oligonucleotides.
Tighe
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                                                                                                             09-OCT-1998;
07-OCT-1999;
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                                                                                                                                                                                                                                                                                                      WO200021556-A1
                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                           gp120; human
development;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunostimulatory sequence; ISS; immunomodulator; glycoprotein 120;
gp120; human immunodeficiency virus; HIV; immune response; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory oligonucleotides regulate the immune contain an immune-stimulating octanucleotide sequence; cancer, allergic and infectious diseases
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                                                        (DYNA-) DYNAVAX TECHNOLOGIES CORP.
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                                                                                                          98US-0103733.
99US-0415186.
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  Schwartz D,
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Pred. No.
  Takabayashi
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19;
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RESULT 66
RAS1466/c
ID AAS146
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XX 10-MAR
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising a human immunodeficiency virus (HIV) antigen, and an immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS). This sequence represents an ISS that can be used in the composition. An immunostimulatory composition which comprises a gp120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-HIV composition comprises immunostimulatory polynucleotides and HIV glycoprotein gp120 useful for modulating, stimulating an immune response against HIV in an HIV infected individual
                                                                                     Suppressing a respiratory syncytial virus infection by administering immunostimulatory sequence at the site of infection is useful to prevent and treat lower respiratory tract viral infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Respiratory syncytial virus. Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                             10-MAR-2000;
09-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adenovirus; measles virus; mumps virus; parainfluenza virus; rubella virus; poxvirus; parvovirus; hantavirus; varicella v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS14666;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 6 A; 8 C; 3 G; 6 T; 0 other;
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                                                                                                                                                                                                               WPI; 2001-607438/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-2001; 2001WO-US07839.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory sequence; ISS; ds; antiviral; immunogen;
respiratory syncytial virus; RSV; influenza virus; rhinovirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunostimulatory sequence, ISS #3.
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2001US-0802686.
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Pred. No.
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Disclosure; Page 15; 40pp; English

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RESULT 67
AAH75994/c
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microcarrier complexes. The complexes comprise an immunostimulatory sequence (ISS), e.g. the present sequence, linked to a nonbiodegradable microcarrier provided that if the microcarrier is gold, latex or magnetic then the linkage is not biotin/avidin. The complex is useful for modulating an immune response (especially stimulating a Thi-type response or suppressing a Th2-type response), increasing interferon-gamma (especially in a patient suffering from idiopathic pulmonary fibrosis), increasing interferon-alpha (especially in patients suffering from viral increasing interferon-alpha (especially in patients suffering from viral increasing interferon-alpha (especially in patients suffering from viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                including influenza virus, rhinovirus, adenovirus, measles virus, mumps virus, parainfluenza virus, rubella virus, poxvirus, parvovirus, hantavirus and varicella virus. A kit for carrying out the administration is also included. Unlike the prior art antiviral agent ribavirin, which is a potential teratogen, the invention provides a treatment which does not carry unacceptable side effects. Other prior art medicaments treat the symptoms only, whilst the invention treats the
                                                                                                                                                                                                                                The present invention relates to immunomodulatory polynucleotide/
                                                                                                                                                                                                                                                                                                                                                                                                Immunomodulatory polynucleotide/microcarrier complexes comprise an immunostimulatory sequence containing polynucleotide linked to a nonbiodegradable microcarrier
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09-MAR-2001; 2001US-0802376.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
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                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Nest G,
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Th2-type immune response; interferon; idiopathic pulmonary fibrosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection. The present sequence is an ISS of the invention.
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                                                                                                                                                                                                                                                                                                                                 Page 18; 61pp; English.
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Pred. No.
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GAACGTTCGAGATGA 20

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RESULT 68
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Matches
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Best Local Similarity
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Best Local
                                                                                                (IIS) of conjugate molecules (CMS) comprising immunostimulatory sequences (ISS) of polynucleotides and antigens. The exent of conjugation affects the immunological properties (e.g. the extent of antigen-specific antibody formation, including Th1-associated antibody formation) so the conjugates are used for altering the type and extent of immune response. (I) and (II) have immunomodulatory, immunosuppressive and antiallergic activities, and can be used in the modulation of immune responses via the stimulation of Th1 lymphocytes and Th1-associated cytokines, and suppression of Th2 lymphocytes and Th1-associated cytokines, and (II) of conjugate molecules may be used for modulating immune responses in individuals e.g. for the treatment of an allergic condition. (I) and (II) may be used to modulate immune responses and therefore prevent potentially harmful reactions to antigens. The present sequence
                                                                                                                                                                                                                                                                                                                                                                     Populations of conjugate molecules comprising polynucleotide immunostimulatory sequences polynucleotides and antigens, useful for controlling immune responses -
                                                         Sequence 23 BP; 6 A; 8 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-NOV-1999; 99US-0165467.
14-NOV-2000; 2000US-0713136.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory sequence; ISS; immunomodulatory; immune response; antigen; antiallergic; modulation; Th1 lymphocyte stimulation; alle Th1-associated cytokine; Th2 lymphocyte suppression; cytokine; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulatory sequence (ISS) SEQ ID NO:3
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15;
                                                                                     present invention.
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Nest G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; DNA; 23
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Conservative
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100.0%; P
tive 0;
            71.4%; Score 15; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                         97pp; English.
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Pred. No.
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                          Length 23;
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AAS16339/c

ID AAS16339 standard; DNA; 23

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AC AAS16339;

XX

DT 14-FEB-2002 (first entry)

XX

DE ISS polynucleotide #3 usef

XX

Herpes simplex virus; HSV

XW Herpes simplex virus; HSV

XW HSV-1; HSV-2; chicken pox;

XX

XX

Virucide; ss.

XX

Synthetic.

XX

PN W0200168103-A2.
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                                                                                                                                                                                                                                    RESULT 70
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                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modulating immune response to a second antigen in humans involves administering an immunostimulatory polynucleotide comprising an immunostimulatory sequence and a first antigen -
                                                                                                                                                                                                                                                                                                                                                                                                   a second antigen in an individual, involving administering to the individual an immunomodulatory polynucleotide comprising an immunostimulatory sequence (ISS) and a first antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2001
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AAF77042 standard; DNA;
                                                              Herpes simplex virus; HSV infection; immunostimulatory sequence; ISS; immune response; alphaherpesvirinae; herpes virus zoster virus; VZV; HSV-1; HSV-2; chicken pox; herpes labialis; cold sore; genital herpes; virucide; ss.
                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 6 A; 8 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to modulating an immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 15; 63pp; English.
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                                                                                                                               polynucleotide #3 useful for treating herpes virus infections
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immune; antigen; immunostimulatory; ds
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                                                                                                                                                        (first entry)
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                                                                                                                                                                                                            23 BP
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AAS16350/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to novel methods of treating, preventing, or reducing the severity or recurrence of a symptom of herpes simplex virus (HSV) infection in an individual who has been exposed to or who is infected with HSV. The method comprises administering a polynuclectide having an immune response. A composition containing ISS is administered induces an immune response. A composition containing ISS is administered vithout a HSV (alphaherpesvirinae) antigen. The composition can be concluded in a kit for ameliorating or preventing a symptom of HSV infection caused by herpes virus zoster virus (VZV), HSV-1 and particularly HSV-2. Such HSV infections include chicken pox, herpes cone of the ISS polynucleotides of the invention.

Note: The present sequence is shown as single stranded in the specification, but the patentees state on page 20 that this sequence may be desification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          preventing, reducing the severity or reducing the recurrence of an infection or symptom of herpes simplex virus (HSV), e.g. HSV-2, comprises administering an immunostimulatory sequence to an individual
                                                                                                                                                                                   Animal papillomavirus infection; human papillomavirus; HPV; STD; wart; sexually transmitted disease; cervical cancer; immune response; immunostimulatory sequence; ISS; virucide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-MAR-2000;
09-MAR-2001;
                            10-MAR-2000;
09-MAR-2001;
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                                                                                                  20-SEP-2001.
                                                                                                                               WO200168117-A2
                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                             ISS polynucleotide #3 useful for treating papillomavirus infections.
                                                                                                                                                                                                                                                                           14-FEB-2002
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(DYNA-) DYNAVAX TECHNOLOGIES
                                                                      12-MAR-2001; 2001WO-US07842
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2001US-0802518.
                            2000US-188265P.
2001US-0802445.
                                                                                                                                                                                                                                                                           (first entry)
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Best Local S
Matches 15
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Note: The present sequence is shown as single stranded in the specification, but the patentees state on page 20 that this sequence may
                                                                                                                                                                                                                                                                                     Cytotoxic T lymphocyte; CTL; T cell; tumour load; cancer radiotherapy; immunostimulatory sequence oligonucleotide; ISS-ODN; chemotherapy; immunosuppression; transplantation; and colimbure disease; infection; acquired immune deficiency syndrome; AIDS; intracellular pathogen; cytomegalovirus; mycobacterial infection; Epstein-Barr virus; varicella zoster virus; human immunodeficiency virus; HIV;
               04-OCT-2001
                                             WO200172123-A1
                                                                                                                            misc_feature
                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                         phosphorothicate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 GAACGTTCGAGATGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunostimulatory sequence to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ng the severity or recurrence of a symptom of papillomavirus in an individual that has been exposed to or who is infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6 A; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates to novel methods of treating,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                     /*tag= a
/mod_base= OTHER
                                                                                                                                           note= "Phosphorothioate backbone"
                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                    sequence oligonucleotide (ISS-ODN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71.4%; Score 15; 100.0%; Pred. No.
                                                                           "This residue is
in page 44 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                         88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising a polynucleotide to the individual -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 24;
                                                                           not found in the sequence specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preventing,
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RESULT 73
ABA03835/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       individual, comprising administering an immunostimulatory sequence coligonuclectide (ISS-ODN). The immunostimulatory nucleic acids of the cinvention are used in CD4+ T cell-deficient individuals to decrease tumour load, to treat a primary or acquired immunodeficiency, and due to cancer radiotherapy or chemotherapy or immunosuppression following to cancer radiotherapy or chemotherapy or immunosuppression following comparison to transplantation, or autoimmune disease treatment, or is acquired immunodeficiency syndrome (AIDS). The nucleic acids may be used to treat a person at risk of becoming CD4+ T cell-deficient, cor is acquired infection, particularly by an intracellular pathogen, consed to treat infection, particularly by an intracellular pathogen, consed to treat infection, particularly by an intracellular pathogen, consed to treat infection, particularly by an intracellular pathogen, consed to treat a person at risk of becoming CD4+ T cell-deficient, cor used to treat infection, particularly by an intracellular pathogen, consed to treat infection, particularly by an intracellular pathogen, consectively probacterium tuberculosis, consectively pathogen, consectively pathogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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11-MAY-2000;
05-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REGC )
                                                                                                                                                                                                          immunomodulation; immunostimulation; phosphorothicate; immunomodulator; antiallergic; antibacterial; antiprotozoal; antiparasitic; hepatotropic; nephrotropic; interferon-alpha stimulator; interferon-gamma stimulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Increasing antigen-specific cytotoxic T lymphocyte activity in a CD4+cell deficient individual, useful to treat immunodeficiency and block HIV infection, comprises administering immunostimulatory nucleic acid
                                                                                                               immunoglobulin E stimulator; immune response; IPP; scleroderma; malaria; idiopathic pulmonary fibrosis; cutaneous radiation-induced fibrosis; hepatic fibrosis; renal fibrosis; infectious disease; leishmaniasis; mycobacterial disease; toxoplasmosis; schistosomiasis; chionorchiasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA03835 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for increasing antigen-specific cytotoxic T lymphocyte (CTL) activity in a CD4+ T cell-deficient
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-MAR-2001; 2001WO-US10118.
                                                                                        allergy; allergy-induced
                                                                                                                                                                                                                                                            Immunomodulatory polynucleotide/microcarrier complex; IMP/MC; immunomodulation; immunostimulation; phosphorothicate; immunom
                                                                                                                                                                                                                                                                                                                                     Immunostimulatory sequence (ISS) SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                          12-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 GAACGTTCGAGATGA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 53; 91pp; English.
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; 2000US-203567P.
; 2000US-215895P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VETERANS
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                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Richman DD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 C; 7 G; 6 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AFFAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22
                                                                                          asthma; prophylactic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Horner AA
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19;
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WO200168144-A2

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ABA03846/c
ID ABA038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microcarrier (IMP/MC) complex (I), which comprises a polynucleotide chaving an immunostimulatory sequence (ISS) linked to a biodegradable microcarrier (MC). The ISS comprises the sequence: 5, CG-3, where the CC microcarrier (MC). The ISS comprises the sequence: 5, CG-3, where the CC antiallergic, antibacterial, antiprotozoal, antiparasitic, hepatotropic cativities. It can be used as an interferon (IFN)-alpha cCC antiallergic, antibacterial, antiprotozoal, antiparasitic, hepatotropic cCC (I) can be used for modulating the immune response of individuals creating idiopathic pulmonary fibrosis (IFP), scleroderma, cutaneous particularly humans. The IMP/MC complex is particularly useful for treating idiopathic pulmonary fibrosis (IFP), scleroderma, cutaneous cradiation-induced fibrosis, hepatic fibrosis including schistosomiasis-induced hepatic fibrosis, renal fibrosis, infectious cradiation-induced by cellular pathogen (e.g. a mycobacterial disease, complexed by cellular pathogen (e.g. a mycobacterial disease, complexed complexed by cellular pathogen (e.g. a mycobacterial disease, complexed by cellular p
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Best Local S
Matches 15
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09-MAR-2001;
                                     20-SEP-2001
                                                                                                       WO200168077-A2
                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABA03846;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABA03846 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New immunomodulatory polynucleotide/microcarrier complex, useful for modulating the immune response of individuals, particularly humans, of treating idiopathic pulmonary fibrosis, scleroderma, malaria or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Van Nest G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAR-2001; 2001WO-US07848
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                                                                                                                                                                                                                                                                                                                                   Immunostimulatory sequence
                                                                                                                                                                                                                                                                                                                                                                                                       12-FEB-2002
                                                                                                                                                                                                                                      immunomodulation;
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                                                                                                                                                                                                                                                                 mmunostimulatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAACGTTCGAGATGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71.4%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 18; 63pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 8 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-188303P.
2001US-0802359.
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                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                  sequence;
virucide;
                                                                                                                                                                                                                                                                                                                                   (ISS)
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%; Pred. No. 19;
0; Mismatches
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                                                                                                                                                                                                                               ISS;
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                                                                                                                                                                                                                           immunostimulation;
therapy; viraemia;
                                                                                                                                                                                                                                                                                                                                   SEQ ID NO:3.
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                                                                                                                                                                                                                                  viral infection; ss.
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laria or
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RESULT 75
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Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a method for reducing severity of a symptom of virus infection in an individual infected with a virus. The method comprises administering a composition consisting of a polynucleotide having an immunostimulatory sequence (ISS). The ISS comprises the sequence 5'-C,G,pyrimidine,pyrimidine,C,G-3'. An antigen is administered in conjunction with the composition. ISS has virucide activity and can be used in gene therapy. The method using the ISS can be used for suppressing, ameliorating and/or preventing viral infections to an individual who may be at risk of being exposed to, exposed to or infected by a virus. It may also be used in reducing the recurrence or duration of a symptom of viral infection, delaying the development of a virus infection, and reducing viraemia or blood levels of virus antigens. The present sequence represents an ISS given in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-MAR-2000;
09-MAR-2001;
                                                          10-MAR-2000;
09-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                ABA03858;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprises administering a polynucleotide having an immunostimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reducing severity, recurrence or duration of symptom of virus infection, or reducing viremia or blood levels of virus antigen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-048999/06
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                                                                                                          12-MAR-2001;
                                                                                                                                         20-SEP-2001.
                                                                                                                                                                        WO200168078-A2
                                                                                                                                                                                                                                    fatigue;
                                                                                                                                                                                                                                                   hepatitis B virus; hepatitis C virus; virucide; anti-inflammatory; hepatotropic; gene therapy; hepatitis infection; viraemia; jaundice;
                                                                                                                                                                                                                                                                                  Immunostimulatory sequence; ISS; immunomodulation; HBV; HCV; infection;
                                                                                                                                                                                                                                                                                                               Immunostimulatory sequence (ISS) SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                            ABA03858 standard; DNA;
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                             (DYNA-) DYNAVAX TECHNOLOGIES CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAACGTTCGAGATGA 20
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llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-188302P.
2001US-0802685.
                                                          2000US-188301P.
2001US-0802370.
                                                                                                          2001WO-US07931.
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Pred. No.
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Job time : 150 secs
                                                                                                                                                                                                                                                                                                         The present invention describes a method for reducing viraemia or blood CC levels of a hepatitis virus antigen in an individual infected with CC hepatitis B virus (HBV). The method comprises administering a composition CC comprising a polynucleotide having an immunostimulatory sequence (ISS) CC to the individual, where the ISS comprises the sequence 5'-C, G-3', and CB WB antigen is not administered in conjunction with administration of the composition, and where the composition is administered in an amount CC antigen. ISS has virucide, anti-inflammatory and hepatotropic activities, CC and can be used in gene therapy. The method can be used for suppressing CC and/or ameliorating hepatitis infection in an individual, especially for preventing, palliating, ameliorating, reducing and/or eliminating one or CC more symptoms of HBV or HCV (hepatitis C virus) infection without CC administering HBV or HCV (hepatitis viral antigen in blood. ISS-containing CC polynucleotides may also be used to improve physical symptoms such as GC indings associated with hepatitis such as blood levels of liver enzymes, portal hypertension, or cirrhosis. The present sequence represents an CC invention.
                                                                                                                                                                     Matches
                                                                                                                                                                                     Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reducing viremia and blood levels of hepatitis virus antigen in an individual infected with hepatitis B virus, comprises administering a composition comprising a polynucleotide having an immunostimulatory sequence
                                                                                                                                                                                                                                                        Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 20; 43pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-049000/06.
                                                                                6 GAACGTTCGAGATGA 20
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15 GAACGTTCGAGATGA 1
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15; Conserva
                                                                                                                                                                 71.4%; Score 15; DB 24; llarity 100.0%; Pred. No. 19; Conservative 0; Mismatches 0
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1: /cgn2_6/ptodata/1/pubpna/US07 PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US07 NEW PUB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US07 NEW PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08 NEW PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08 NEW PUB.seq:*

8: /cgn2_6/ptodata/1/pubpna/US08 PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US09 PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US09 PUBCOMB.seq:*

10: /cgn2_6/ptodata/1/pubpna/US09 PUBCOMB.seq:*

11: /cgn2_6/ptodata/1/pubpna/US09 NEW PUB.seq:*

13: /cgn2_6/ptodata/1/pubpna/US09 NEW PUB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US10A PUBCOMB.seq:*

15: /cgn2_6/ptodata/1/pubpna/US10A PUBCOMB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10A PUBCOMB.seq:*

17: /cgn2_6/ptodata/1/pubpna/US10B NEW PUB.seq:*

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17: /cgn2_6/ptodata/1/pubpna/US10B NEW PUB.seq:*

18: /cgn2_6/ptodata/1/pubpna/US10B NEW PUB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10B NEW PUB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10B NEW PUB.seq:*
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21
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    satisfying
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US-10-03-243-132

US-09-927-422A-16

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US-10-177-826-41

US-10-13-243-19

US-10-177-826-52

US-10-177-826-52

US-10-177-826-52

US-10-177-826-36

US-10-177-823-36

US-10-177-823-36

US-10-177-826-31

US-10-177-826-31

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  sequence
1132, App
116, Appl
111, Appl
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    US-09-802-686-1
US-09-802-685-2
US-09-802-685-2
US-09-802-685-1
US-09-802-376-1
US-09-802-376-2
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US-09-902-376-2
US-09-902-376-2
US-09-902-376-2
US-09-902-445-1
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APPLICANT: FEARON, Karen L.

APPLICANT: DINA, Dino

TITLE OF INVENTION: METHODS OF USING THE SAME

TITLE OF INVENTION: METHODS OF USING THE SAME

FILE REFERENCE: 377882001800

CURRENT APPLICATION NUMBER: US/10/033,243

CURRENT FILING DATE: 2002-04-03

PRIOR APPLICATION NUMBER: 60/258,675

PRIOR APPLICATION NUMBER: 60/258,675

PRIOR APPLICATION NUMBER: 30/258,675

PRIOR PILING DATE: 2000-12-27

INUMBER OF SEO ID NOS: 133

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 132

LENGTH: 21

TYPE: DNA

ORGANISM: Artificial Sequence
                  CURRENT APPLICATION NUMBER: US/09/927,422A
CURRENT FILLING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: U.S. 09/802,359
PRIOR FILLING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: U.S. 60/188,30
PRIOR FILLING DATE: 2000-03-10
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-927-422A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1
US-10-033-243-132
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 16, Application US/09927422A Publication No. US20030022852A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 132, Application No. US20 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                              APPLICANT: Fearon, Karen L.
APPLICANT: Dina, Dino
TITLE OF INVENTION: BIODEGRADABLE IMMUNOMODULATORY
TITLE OF INVENTION: FORMULATIONS AND METHODS FOR U
FILE REFERENCE: 377882001420
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Van Nest, Gary
APPLICANT: Tuck, Stephen
APPLICANT: Fearon, Karen L.
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TYPE: DNA
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US-10-176-883-104

US-10-177-826-98

US-10-177-826-104

US-10-177-826-104

US-10-176-883-33

US-10-176-883-83

US-10-176-883-83

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US-10-176-883-83
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CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR PILING DATE: 2002-04-23
NUMBER OF ESO ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 41
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
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US-10-177-826-41
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 41, Application US/10177826
Publication No. US20030199466A1
GENERAL INFORMATION:
                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 41
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Best Local Similarity
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Best Local Similarity
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                                                                                                               CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Fearon, Karen APPLICANT: Dina, Dino APPLICANT: Tuck, Stephen
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APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
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                       LENGTH: 19
TYPE: DNA
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ORGANISM: Artificial Sequence
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CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
PRIOR FILING DATE: 2002-04-23
PRIOR FILING DATE: 141
SOFTWARE: FABESEQ for Windows Version 4.0
SEQ ID NO 52
ENGTH: 22
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US-10-033-243-19
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SEQ ID NO 19
LENCTH: 19
TYPE: DNA
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Best Local Similarity
Matches 19; Conserv
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CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
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PRIOR FILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 133
                                                                                                                                                                                                                                                      TITLE OF INVENTION: CHIMERIC IMMUNOMOBULATORY COMPOUNDS AND TITLE OF INVENTION: METHODS OF USING THE SAME-I FILE REFERENCE: 377882002000
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APPLICANT: DINA, Dino
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OTHER INFORMATION: Synthetic construct
                                  ORGANISM: Artificial Sequence
                                                       TYPE: DNA
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Tuck, Stephen
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100.0%; Pred. No.
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                                                                        US-10-033-243-30
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LENGTH: 22
TYPE: DNA
                                                                                                                                                               NUMBER OF SEQ ID NOS: 133
SOFTWARE: FRANSEQ for Windows Version 4.0
SEQ ID NO 30
LENGTH: 22
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Best Local Similarity
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Best Local Similarity Matches 19; Conserv
                                    Query Match
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CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR PLING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-12-27
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APPLICANT: Dina, I
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                                                                                                                                                 TYPE: DNA
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90.5%; Score 19; DB 15; Length 22; 100.0%; Pred. No. 0.064; tive 0; Mismatches 0; Indels
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TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND TITLE OF INVENTION: METHODS OF USING THE SAME-I FILE REFERENCE: 377882002000

CURRENT APPLICATION NUMBER: US/10/176,883

CURRENT FILING DATE: 2002-06-21

PRIOR APPLICATION NUMBER: 60/299,883

PRIOR FILING DATE: 2001-06-21

PRIOR APPLICATION NUMBER: 60/375,253

PRIOR FILING DATE: 2002-04-23

NUMBER OF SEQ ID NOS: 141

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 36
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US-10-176-883-36
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; OTHER INFORMATION: Synthetic construct
US-10-176-883-36
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                                                                                                    ; OTHER INFORMATION: Synthetic construct US-10-177-826-36
                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-66-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 36
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Publication No.
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Publication No.
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Best Local Similarity
                                               Query Match
Best Local Similarity
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                                   Matches
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CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
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                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                    FEATURE:
                                                                                                                                                                                   ENGTH: 18
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                                   16; Conservative
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TCGAACGTTCGAGATG 19
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Tuck, Stephen
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No. US20030199466A1
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                                               76.2%; Score 16; 100.0%; Pred. No.
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                                 Mismatches
                                                                 DB 13; Length 18;
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Best Local S
Matches 16
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Matches
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APPLICANT: Dina, I
APPLICANT: Tuck, S
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                                                                                                                  Local Similarity
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US-10-033-243-14

US-10-033-243-14

; Sequence 14, Application US/10033243

; Publication No. US20030049266A1
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                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 139
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APPLICANT: FEARON, Karen L.
APPLICANT: DINA, Dino
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APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
CURRENT APPLICATION UNDBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR PILING DATE: 2001-06-21
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CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR PILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 133
                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                             GAACGTTCGAGATGAT 21
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GAACGTTCGAGATGAT 23
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                                                                                                76.2%; Score 16; DB llarity 100.0%; Pred. No. 3. Conservative 0; Mismatches
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RESULT 13 US-10-177-826-139

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                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: us 60/262321
PRIOR FILING DATE: 2001-01-17
PRIOR APPLICATION NUMBER: us 60/202,274
PRIOR FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 20
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Best Local Similarity
Matches 16; Conserv
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Best Local 9
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                                                                             Matches
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APPLICANT: Takabayashi, Kenji
TITLE OF INVENTION: Agents that Modulate DNA-PK Activity and
TITLE OF INVENTION: Methods of Use Thereof
FILE REFERENCE: 06510168US1
CURRENT APPLICATION NUMBER: US/09/848,986
CURRENT FILING DATE: 2001-05-03
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PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTMARE: FASTSEQ for Windows Version 4.0
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                Similarity
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GAACGTTCGAGATGA 20
                                  GAACGTTCGAGATGA 20
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Tuck, Stephen
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                                                                           Conservative
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                                                                       71.4%; Score 15;
100.0%; Pred. No.
tive 0; Mismatc
                                                                       ore 15; DB 13; Length 20; red. No. 16; Mismatches 0; Indels
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RESULT 15

RESULT 17 US-09-802-686-2

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US-10-233-121A-21
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CURRENT FILING DATE: 2003-03-13-11
PRIOR APPLICATION NUMBER: US 09/848,986
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/202,274
PRIOR APPLICATION NUMBER: US 60/262,321
PRIOR APPLICATION NUMBER: US 60/262,321
PRIOR APPLICATION NUMBER: US 60/262,321
PRIOR FILING DATE: 2001-01-17
NUMBER OF SEQ ID NOS: 21
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SEQ ID NO 1
LENGTH: 22
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SEQ ID NO 21
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                   71.4%; Score 15; DB 9; Length 22; Best Local Similarity 100.0%; Pred. No. 16; Matches 15; Conservative
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Best Local Similarity
Matches 15; Conserv
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Patent No. US20010046967A1
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TITLE OF INVENTION: METHODS OF PREVENTING AND TREATING
TITLE OF INVENTION: RESPIRATORY VIRAL INFECTION USING IMMUNOMODULATORY
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCES
FILE REFERENCE: 377882000900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dynavax Technologies Corporation APPLICANT: Van Nest, Gary
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CURRENT FILING DATE: 2001-03-09
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APPLICANT: LOIS, AUGUST
APPLICANT: TAKABAYASHI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: TAKABAYASHI, KENJI
TITLE OF INVENTION: AGENTS THAT MODULATE DNA-PK ACTIVITY AND
TITLE OF INVENTION: METHODS OF USE THEREOF
                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: UCAL-168DIV
                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                    OTHER INFORMATION: Polynucleotide containing
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US-09-802-685-2
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US-09-802-685-1
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SEQ ID NO 2
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
GENERAL INFORMATION:
APPLICANT: Van Nest, Gary
APPLICANT: Eiden, Joseph J., Jr.
TITLE OF INVENTION: METHODS OF PREVENTING AND TREATING VIRAL
                                                                          Sequence 2, Application US/09802685
Patent No. US20020028784A1
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Best Local S
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Patent No. US20020028784A1
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CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/188,583
PRIOR FILING DATE: 2000-03-10
NUMBER OF SEQ ID NOS: 10
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TITLE OF INVENTION: METHODS OF PREVENTING AND TREATING
TITLE OF INVENTION: RESPIRATORY VIRAL INFECTION USING
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCES
FILE REFERENCE: 377882000900
FILE REFERENCE: 377882000900
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APPLICANT: Eiden, Joseph J., Jr.
TITLE OF INVENTION: METHODS OF PREVENTING AND TREATING VIRAL
TITLE OF INVENTION: INFECTIONS USING IMMUNOMODULATORY POLYNUCLEOTIDE SEQUENCES
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CURRENT APPLICATION NUMBER: US/09/802,685
CURRENT FILING DATE: 2001-03-09
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                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Polynucleotide containing
                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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US-09-802-376-1
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US-09-791-500-1
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APPLICANT: Raz, Eyal:
APPLICANT: Raz, Eyal:
APPLICANT: Raz, Eyal:
APPLICANT: Raz, Eyal:
APPLICANT: Razh Eval:
APPLICANTION: Method for Treating Inflammatory Bowel
TITLE OF INVENTION: Method for Treating Inflammatory.
FILE OF INVENTION: MUMBER: US/09/791,500
CURRENT APPLICATION NUMBER: US/09/791,500
CURRENT FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 39
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
TYDER. DNA
CURRENT APPLICATION NUMBER: US/09/802,376
CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/188,557
PRIOR FILING DATE: 2000-03-10
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
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Patent No. US20020055477A1
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/802,685
CURRENT FILING DATE: 2001-03-09
                                                                                                                                                            APPLICANT: Van Nest, Gary
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: IMMUNOMODULATORY FORMULATIONS AND METHODS FOR USE THEREOF
FILE REFERENCE: 37788201700
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; FEATURE:
; OTHER INFORMATION:
US-09-802-370-1
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; ORANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Polynucleotide containing
US-09-802-376-1
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US-09-802-376-2
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                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 2
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Matches 15; Conserv
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Best Local (
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APPLICANT: Eiden, JOSEPh J. Jr.
APPLICANT: Eiden, JOSEPh J. Jr.
TITLE OF INVENTION: METHODS OP SUPPRESSING HEPATITIS VIRUS
TITLE OF INVENTION: INFECTION USING IMMUNOMODULATORY POLYN
FILE REFERENCE: 377882001200
CURRENT APPLICATION NUMBER: US/09/802,370
CURRENT FILING DATE: 2001-09-24 (
PRIOR APPLICATION NUMBER: 60/188,301
PRIOR APPLICATION NUMBER: 60/188,301
PRIOR FILING DATE: 2000-03-10
NUMBER OF SEQ ID NOS: 8
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CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/188,557
PRIOR FILING DATE: 2000-03-10
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APPLICANT: Tuck, Stephen
TITLE OF INVENTION: IMMUNOMODULATORY FORMULATIONS AND METHODS FOR USE THEREOF
FILE REFERENCE: 37788201700
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ORGANISM: Artificial Sequence
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                                                                      Polynucleotide containing
71.4%;
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Pred.
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US-09-802-445-1
; Sequence 1, Application US/09802445
; Patent No. US20020107212A1
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US-09-802-370-2
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Matches
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APPLICANT: Van Nest, Gary
APPLICANT: Eiden, Joseph J. Jr.
APPLICANT: Eiden, JOSEPH J. Jr.
TITLE OF INVENTION: METHODS OF REDUCING PAPILLOMAVIRUS INFECTION USING IMMUNOMODULATO.
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCES
FILE REFERENCE: 377882001300
                                                                                                                                                                                                                              SEQ ID NO 1
LENGTH: 22
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: 60/188,301
PRIOR FILING DATE: 2000-03-10
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APPLICANT: Eiden, Joseph J. Jr.
TITLE OF INVENTION: METHODS OF SUPPRESSING HEPATITIS VIRUS
TITLE OF INVENTION: INFECTION USING IMMUNOMODULATORY POLYNUCLEOTIDE SEQUENCES
                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/188,265 PRIOR FILING DATE: 2000-03-10
                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/802,445
CURRENT FILING DATE: 2001-09-24
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                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
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GAACGTTCGAGATGA 22
                                 GAACGTTCGAGATGA 20
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US-09-820-484-1
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SEQ ID NO 1
LENGTH: 22
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Patent No. US20020142977A1
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Patent No. US20020107212A1
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APPLICANT: Eiden, Joseph J. Jr.
TITLE OF INVENTION: METHODS OF REDUCING PAPILLOMAVIRUS INFECTION USING IMMUNOMODULATC
TITLE OF INVENTION: POLYNUCLECTIDE SEQUENCES
FILE REFERENCE: 377882001300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cho, Hearn Jay
APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Method for Increasing a Cytotoxic T
TITLE OF INVENTION: Lymphocyte Response in vivo.
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SOFTWARE: FastSEQ for Windows Version 4.0
EQ ID NO 2
LENGTH: 22
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CURRENT APPLICATION NUMBER: US/09/820,484
CURRENT FILING DATE: 2001-03-28
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CURRENT FILING DATE: 2001-09-24
                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                           OTHER INFORMATION: Disulfide-linked phosphorothicate ISS-ODN NAME/KEY: modified base LOCATION: (1)...(1)
OTHER INFORMATION: disulfide thymine
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                                                                     Local Similarity 100.
les 15; Conservative
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                        GAACGTTCGAGATGA 20
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US-09-820-484-3
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Patent No. US20020142978A1
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Best Local Similarity
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; OTHER INFORMATION: phosphorothicate ISS-ODN US-09-820-484-3
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LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Patent No. US20020142977A1
GENERAL INFORMATION:
                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                          APPLICANT: Nguyen, Minh-Duc
TITLE OF INVENTION: Synergistic Improvements
TITLE OF INVENTION: Polynucleotide Vaccines
FILE REFERENCE: 6510-203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/828,505
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: 60/195,890
PRIOR FILING DATE: 2000-04-07
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TITLE OF INVENTION: Method for Increasing a
TITLE OF INVENTION: Lymphocyte Response in
FILE REFERENCE: 06510-188US1
                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                     OTHER INFORMATION: Immunomodulatory nucleic acid sequence
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71.4%; 5--
100.0%; Pr
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 22
TYPE: DNA
ORGANIEM: Artificial sequence
FEATURE: INFORMATION: Oligodeoxynucleotide
US-09-967-881-2
RESULT 32
US-09-927-422A-2
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US-09-927-422A-1
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US-09-967-881-2
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                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
TYPE: DNA
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Best Local Similarity
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APPLICANT: Institut National de la Sante et de la Recherche M
APPLICANT: Carpentier, Antoine
TITLE OF INVENTION: Use of Stabilised Oligonucleotides for Preparing A Medicament wit
TITLE OF INVENTION: Antitumor Activity
FILE REFERENCE: 267/246 US
CURRENT APPLICATION NUMBER: US/09/967,881
CURRENT FILING DATE: 2001-09-28
RUMBER OF SEQ ID NOS: 48
                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/927,422A CURRENT FILING DATE: 2001-08-10 PRIOR APPLICATION NUMBER: U.S. 09/802,359 PRIOR FILING DATE: 2001-03-09 PRIOR APPLICATION NUMBER: U.S. 60/188,30 PRIOR FILING DATE: 2000-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: BIODEGRADABLE IMMUNOMODULATORY TITLE OF INVENTION: FORMULATIONS AND METHODS FOR USE THEREOF FILE REFERENCE: 377882001420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Van Nest, Gary APPLICANT: Tuck, Stephen
                                                                                                                                                                                                                                                             OTHER INFORMATION: Polynucleotide containing CG
                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                 Local
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Fearon, Karen L.
Dina, Dino
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US-09-927-884-1

Sequence 1, Application US/09927884 Publication No. US20030059773A1

GENERAL INFORMATION:

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CURRENT APPLICATION NUMBER: US/09/738,046A
CURRENT FILING DATE: 2000-12-15
NUMBER OF SEQ ID NOS: 3
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
SEQ ID NO 3
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: INFORMATION: artificial sequence containing CpG sequence
US-09-738-046A-3
RESULT 34
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US-09-738-046A-3
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SEQ ID NO 2
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: FELCNER, PHILIP L.
APPLICANT: ZELPHATI, OLIVIER
TITLE OF INVENTION: INTRACELLULAR PROTEIN DELIVERY
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF US
FILE REFERENCE: GTSYS.004A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09927422A Publication No. US20030022852A1
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                                                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/927,422A CURRENT FILING DATE: 2001-08-10 PRIOR APPLICATION NUMBER: U.S. 09/802,359 PRIOR FILING DATE: 2001-03-09 PRIOR APPLICATION NUMBER: U.S. 60/188,30 PRIOR FILING DATE: 2000-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Fearon, Karen L.
APPLICANT: Dina, Dino,
TITLE OF INVENTION: BIODEGRADABLE IMMUNOMODULATORY
TITLE OF INVENTION: FORMULATIONS AND METHODS FOR USE THEREOF
FILE REFERENCE: 377882001420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Van Nest, Gary
APPLICANT: Tuck, Stephen
APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
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                                                       GAACGTTCGAGATGA 22
                                                                                                                              Conservative
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                                                                                                                              71.4%; Score 15;
100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                  Mismatches
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                                                                                                                                                                  Length 22
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APPLICANT: DIA, DIAO

APPLICANT: DIAO, DIAO

TITLE OF INVENTION: IMMUNOMODULATORY FORMULATIONS AND

TITLE OF INVENTION: METHODS FOR USE THEREOF

FILE REFERENCE: 377882001720

CURRENT APPLICATION NUMBER: US/09/927,884

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: U.S. 09/802,376

PRIOR APPLICATION NUMBER: U.S. 60/188,557

PRIOR APPLICATION NUMBER: U.S. 60/188,557

PRIOR APPLICATION NUMBER: U.S. 60/188,557

PRIOR APPLICATION NUMBER: U.S. 50/188,557

PRIOR FILING DATE: 2000-03-10

NUMBER OF SEQ ID NOS: 14

SOSTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 2
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CURRENT APPLICATION NUMBER: US/09/927,884

CURRENT FILING DATE: 2001-08-10

PRIOR PILING DATE: 2001-03-09

PRIOR PILING DATE: 2001-03-09

PRIOR PILING DATE: 2000-03-10

PRIOR PILING DATE: 2000-03-10

NUMBER OF SEQ ID NOS: 14

SOFTWARE: FastSEQ for Windows Version 4.0
                                                                             RESULT 36
US-09-802-359-1
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                 Sequence 1, Application US/09802359 Publication No. US20030129251A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                   Query Match
Best Local :
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LENGTH: 22
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APPLICANT: Van Nest, Gary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Van Nest, Gary APPLICANT: Tuck, Stephen
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                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Polynucleotide containing
                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Polynucleotide containing
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                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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Dino
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Dina, Dino
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                                                                                                                                                                                                                                                                 71.4%; Score 15; DB 11; Length 22; 100.0%; Pred. No. 16;
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US-09-967-464-19
; Sequence 19, Application US/09967464
; Publication No. US20030138453A1
; GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/802,359
CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/188,303
PRIOR APPLICATION NUMBER: 60/188,303
PRIOR FILING DATE: 2000-03-10
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 15; Conserv
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Best Local Similarity
                                                                                                                    APPLICANT: O'Hagan, Derek
APPLICANT: Otten, Gillis
APPLICANT: Donnelly, John J.
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CURRENT FILING DATE: 2001-03-09
                                                                                       APPLICANT:
                                                                                                           APPLICANT:
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                                      APPLICANT:
                                                        APPLICANT:
                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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T: Dubensky, Jr., Thomas W.
INVENTION: MICROPARTICLES FOR DELIVERY OF HETEROLOGOUS NUCLEIC ACIDS
SRENCE: PP16269.004
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                                                  Barnett, Susan
Singh, Mamohan
Ulmer, Jeffrey
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                                                                                                         Polo, John M.
                                                                                                                                                                                                                                                                                                                                                                       71.4%; Score 15; ilarity 100.0%; Pred. No Conservative 0; Mismatc
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Pred. No.
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16;
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PRIOR APPLICATION NUMBER: 60/236,105
PRIOR PILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin version 3.1
SEQ ID NO 19
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                   US-10-340-275-1
                                                                                                                                                                                                                                                     RESULT 40
                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
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LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                           Sequence 1, Application US/10340275 Publication No. US20030143213A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/10214799
Publication No. US20030133988A1
GENERAL INFORMATION:
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                                APPLICANT: Cho, Hearn Jay
APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Method for Increasing a Cytotoxic T
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: UCAL-188DIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/335,263
PRIOR FILING DATE: 2001-10-25
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/214,799
CURRENT FILING DATE: 2002-08-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Fearon, Karen L. and Dina, Dino
TITLE OF INVENTION: IMMUNOMODULATORY COMPOSITIONS,
TITLE OF INVENTION: FORMULATIONS, AND METHODS OF USE THEREOF
FILE REFERENCE: 377882003100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-04-11
CURRENT APPLICATION NUMBER: US/10/340,275
                                                                                                                                                     APPLICANT: Raz, Eyal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Artificial sequence is synthesized
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100.0%; Pred. No.
Live 0; Mismatcl
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PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
PRIOR PRIOR PRICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
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                                                                                                                                         OTHER INFORMATION: phosphorothicate ISS-ODN US-10-340-275-3
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US-10-340-275-3
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APPLICANT: Raz, Eyal
APPLICANT: Cho, Hea
APPLICANT: Richman,
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LENGTH: 22
                                                                                     Query Match
Best Local 9
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Best Local :
                                                                    Matches
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CURRENT FILING DATE: 2003-01-10
PRIOR APPLICATION NUMBER: 09/820,484
PRIOR FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR FILING DATE: 2000-03-28
PRIOR FILING DATE: 2000-03-28
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PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cho, Hearn Jay
APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Method for Increasing a Cytotoxic
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: UCAL-188DIV
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PRIOR FILING DATE: 2001-03-28
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OTHER INFORMATION: disulfide thymine
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ORGANISM: Artificial Sequence
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                             6 GAACGTTCGAGATGA 20
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                                                                    Conservative
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100.0%; Pr
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pred. No.
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Best Local S
Matches 15
                                                              SEQ ID NO 3
LENGTH: 22
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                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/339,885
CURRENT FILING DATE: 2003-01-10
PRIOR APPLICATION NUMBER: 09/820,484
PRIOR FILING DATE: 2001-03-8
PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR APPLICATION NUMBER: US 60/192,537
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A
TITLE OF INVENTION: Method for Increasing a Cytotoxic T
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: UCAL-188CON
                                                                                                                      NUMBER OF SEQ ID NOS: 8
                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/215,895 PRIOR FILING DATE: 2000-07-05
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PRIOR APPLICATION NUMBER: US 60/203,567
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PRIOR FILING DATE: 2000-05-11
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                                                                                                           SOFTWARE: FastSEQ
                                                                                                                                                                                              PRIOR FILING DATE: 2000-05-11
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OTHER INFORMATION: disulfide thymine
                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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No. US20030147870A1
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Richman, Douglas
Horner, Anthony A.
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                                                                                                         for Windows Version 4.0
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100.0%; Pred. No.
tive 0; Mismatc
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US-10-176-883-2
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                                                                                  SEQ ID NO 2
LENGTH: 22
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Best Local :
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LENGTH: 22
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                                                                                                                                                                                                                                                        TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND TITLE OF INVENTION: METHODS OF USING THE SAME-I FILE REFERENCE: 377882002000 CURRENT APPLICATION NUMBER: US/10/176,883 CURRENT FILING DATE: 2002-06-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lois, Augusto F.
APPLICANT: Takabayashi, Kenji
TITLE OF INVENTION: Agents that Modulate DNA-PK Activity and
TITLE OF INVENTION: Methods of Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: us 60/202,274
PRIOR FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FASTSEQ for Windows Version 4.0
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CURRENT APPLICATION NUMBER: US/09/848,986
CURRENT FILING DATE: 2001-05-03
                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/299,883 PRIOR FILING DATE: 2001-06-21
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic construct
                      FEATURE:
                                        ORGANISM: Artificial Sequence
                                                                TYPE: DNA
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100.0%; Pred. No.
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US-10-176-883-24
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; OTHER INFORMATION: Synthetic construct US-10-176-883-34
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PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
LENGTH: 22
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Best Local Similarity
                                                                             SEQ ID NO 34
LENGTH: 22
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Publication No.
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Publication No.
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                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
                                                                                                                                                                                                                                                     APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND TITLE OF INVENTION: METHODS OF USING THE SAME-I FILE REFERENCE: 377882002000
                                                                                                                 PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTWARE: FASTSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-06-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Fearon, Karen
                            TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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Tuck, Stephen
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Pred. No.
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RESULT 49
US-10-176-883-79
US-10-176-883-79
; Sequence 79, Application US/10176883
; Publication No. US20030175731A1
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US-10-176-883-44
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Best Local Similarity
Matches 15; Conserv
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                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 79
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SEQ ID NO 44
LENGTH: 22
TYPE: DNA
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                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
                                                                                                                                                                                                                                                  APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR PILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
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                                                                                                                                NUMBER OF SEQ ID NOS: 141
                       TYPE: DNA ORGANISM: Artificial Sequence
OTHER INFORMATION: Synthetic construct
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                 FEATURE:
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100.0%; Pred. No.
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                                     SEQ ID NO 85
LENGTH: 22
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Best Local Similarity
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APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
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APPLICANT: Dina, Dina
APPLICANT: Dina, Dina
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-I
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PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 377882002000
CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
                                                                                                                                             PRIOR APPLICATION NUMBER: 60/299,883 PRIOR FILING DATE: 2001-06-21
                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
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NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                      SOFTWARE:
                                                                                                          PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                              APPLICANT: Fearon, Karen
TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: Synthetic construct
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                                                                          FastSEQ for Windows Version 4.0
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No. US20030175731A1
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Pred. No.
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US-10-176-883-140
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Matches
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Best Local Similarity
Matches 15; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 140 LENGTH: 22
                                                                                                                                                                                                                                                                                                             Sequence 140, Application US/10176883
Publication No. US20030175731A1
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CURRENT APPLICATION NUMBER: US/10/176,883
CURRENT FILING DATE: 2002-06-21
                                                                                                                                            APPLICANT: Dina, Dina
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMMERIC IMMUNOMODULATORY COMPOUNDS
TITLE OF INVENTION: METHODS OF USING THE SAME-I
FILE REFERENCE: 377882002000
CURRENT APPLICATION UNDMERR: US/10/176,883
CURRENT FILING DATE: 2002-06-21
                                                       PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
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                                                                                                        PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
                                                                                                                                                                                                                                                                         APPLICANT: Fearon,
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ORGANISM: Artificial Sequence
FEATURE:
TYPE: DNA
ORGANISM: Artificial Sequence
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Tuck, Stephen
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100.0%; Pr
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US-10-176-883-141
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Publication No. US20030176389A1
GENERAL INFORMATION:
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Best Local :
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Best Local Similarity
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APPLICANT: Raz, Eyal
APPLICANT: Rachmilewitz, Daniel
TITLE OF INVENTION: Method for Treating Inflammatory Bowel
TITLE OF INVENTION: Disease and Other Forms of Gastrointestinal Inflammation.
FILE REFERENCE: UCAL-202CON
CURRENT APPLICATION NUMBER: US/10/412,151
CURRENT FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 09/791,500
PRIOR FILING DATE: 2001-02-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS TITLE OF INVENTION: METHODS OF USING THE SAME-I FILE REFERENCE: 377882002000 CURRENT APPLICATION NUMBER: US/10/176,883 CURRENT FILING DATE: 2002-06-21
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PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEO ID NOS: 141
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TYPE: DNA
ORGANISM: Artificial Sequence
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NAME/KEY: modified_base
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic construct
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Tuck, Stephen
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Pred. No.
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GENERAL INFORMATION:

APPLICANT: Fearon, Karen

APPLICANT: Dina, Dino

APPLICANT: Dina, Dino

APPLICANT: Tuck, Stephen

TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND

TITLE OF INVENTION: METHODS OF USING THE SAME-II

FILE REFERRICE: 377882002001

CURRENT APPLICATION NUMBER: US/10/177,826

CURRENT FILING DATE: 2002-06-21

PRIOR APPLICATION NUMBER: 60/29,883

PRIOR FILING DATE: 2001-06-21

PRIOR APPLICATION NUMBER: 60/375,253

PRIOR APPLICATION NUMBER: 60/375,253

PRIOR APPLICATION NUMBER: 60/375,253
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US-10-177-826-24
| Sequence 24, Application US/10177826
| Publication No. US20030199466A1
| GENERAL INFORMATION:
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US-10-177-826-2
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LENGTH: 22
TYPE: DNA
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Publication No. US20030199466A1
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LENGTH: 22
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                                APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-II
FILE REFERENCE: 377882002001
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
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OTHER INFORMATION:
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OTHER INFORMATION: Synthetic construct
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60/299,883

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RESULT 59
US-10-177-826-44
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SEQ ID NO 34
                                                                                                                                                           Sequence 44, Application US/10177826 Publication No. US20030199466A1 GENERAL INFORMATION:
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Publication No. US20030199466A1
GENERAL INFORMATION:
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Best Local Similarity
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     APPLICANT: Fearon, Karen
APPLICANT: Dina, Dina
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-II
FILE REFERENCE: 379882002001
CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
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NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version
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PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/375,253
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Tuck, Stephen
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100.0%; Pred. No.
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Pred. No.
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16;
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US-10-177-826-84
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US-10-177-826-79
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NUMBER OF SEQ ID NOS: 141

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 79

LENGTH: 22
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                Sequence 84, Application US/10177826
Publication No. US20030199466A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                           Best
APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS AND
TITLE OF INVENTION: METHODS OF USING THE SAME-II
FILE REFERENCE: 377882002001
CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
REFOR APPLICATION NUMBER: 60/299,883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: CHIMERIC IMMUNOMODULATORY COMPOUNDS TITLE OF INVENTION: METHODS OF USING THE SAME-II FILE REFERENCE: 377882002001 CURRENT APPLICATION NUMBER: US/10/177,826 CURRENT FILING DATE: 2002-06-21
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PRIOR APPLICATION NUMBER: 60/-
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
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PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
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14; Conser
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Pred. No. 16;
1; Mismatches
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16;
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RESULT 63
US-10-177-826-134
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PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 85
LENGTH: 22
                                                                         Sequence 134, Application US/10177826 Publication No. US20030199466A1
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Publication No. US20030199466A1
GENERAL INFORMATION:
                                                            GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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APPLICANT: Fearon, Karen
APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
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APPLICANT: Tuck, Stephen
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2002-04-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                   71.4%;
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US-10-177-826-140
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                                                                     Query Match
Best Local S
Matches 15
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SEQ ID NO 140
LENGTH: 22
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Best Local !
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SEQ ID NO 134
LENGTH: 22
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CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2001-06-23
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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APPLICANT: Dina, Dino
APPLICANT: Tuck, Stephen
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CURRENT APPLICATION NUMBER: US/10/177,826
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
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NUMBER OF SEQ ID NOS: 141
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                                                                                                                                                                    OTHER INFORMATION: n = thymine attached to a reactive linking group
                                                                                                                                                                                    FEATURE:
NAME/KEY: modified_base
LOCATION: 1
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                                                                                                                                                                                                                                         OTHER INFORMATION: Synthetic construct
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                                 6 GAACGITCGAGAIGA 20
GAACGITCGAGATGA 22
                                                                         Conservative
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100.0%; Pred. No.
                                                                                         71.4%; Score 15; 100.0%; Pred. No.
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RESULT 65
US-10-177-826-141
; Sequence 141, Application US/10177826
; Publication No. US20030199466A1
; GENERAL INFORMATION:

Fearon, Karen

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RESULT 67
US-10-357-760-2
; Sequence 2, Application US/10357760
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                                                                                                                                                                    Query Match
Best Local S
Matches 15
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 377882001200
CURRENT APPLICATION NUMBER: US/10/357,760
CURRENT FILING DATE: 2003-02-03
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CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: 60/299,883
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/375,253
PRIOR FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Van Nest, Gary
APPLICANT: Eiden, Joseph J. Jr.
TITLE OF INVENTION: INFECTION USING IMMUNOMODULATORY POLYNUCLEOTIDE SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2000-03-10 NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: 60/188,301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US/09/802,370
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                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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Tuck, Stephen
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100.0%; F1
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100.0%; Pr
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                                                                                                 22
                                                                                                                                                                      Score 15; DB; Pred. No. 16; 0; Mismatches
                                                                                                                                                                                       DB 13; Length 22; 16;
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RESULT 69 US-10-033-243-2

Sequence 2, Application US/10033243
Publication No. US20030049266A1
GENERAL INFORMATION:
APPLICANT: FEARON, Karen L.
APPLICANT: DINA, Dino

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Best Local Similarity
Matches 15; Conserv
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CURRENT FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: US 60/264,476
PRIOR FILING DATE: 2001-01-26
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
                                                                        Query Match
Best Local Similarity
Matches 15; Conserv
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LENGTH: 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Van Nest, Gary
APPLICANT: Eiden, Joseph J. Jr.
TITLE OF INVENTION: METHODS OF SUPPRESSING HEPATITIS VIRUS
TITLE OF INVENTION: INFECTION USING IMMUNOMODULATORY POLYI
EILE REFERENCE: 377882001200
                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Method For Treating an HIV-Infected TITLE OF INVENTION: Individual By Combining Immunization With Structured TITLE OF INVENTION: Interruption of Anti-Retroviral Treatment FILE REFERENCE: P-IM 5158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2000-03-10 NUMBER OF SEQ ID NOS: 8
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ORGANISM: Artificial Sequence
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GAACGTTCGAGATGA
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100.0%; Pred. No.
vative 0; Mismatc
                                                                        71.4%; Score 15; DB
100.0%; Pred. No. 16
tive 0; Mismatches
22
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RESULT 71
US-10-033-243-22
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SEQ ID NO 2
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
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Publication No.
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 22
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Best Local Similarity
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Best Local Similarity
APPLICANT: FEARON, KAREN L.
APPLICANT: DINA, DINO
TITLE OF INVENTION: IMMUNOMODULATORY POLYNUCLEOTIDES AND
TITLE OF INVENTION: METHODS OF USING THE SAME
FILE REFERENCE: 377882001800
CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR FILING DATE: 2000-12-27
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CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR FILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 133
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Pred. No.
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US-10-033-243-45
; Sequence 45, Application US/10033243
; Publication No. US20030049266A1
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SEQ ID NO 22
LENGTH: 22
                                                 SOFTWARE: FastSEQ for Windows Version SEQ ID NO 45
LENGTH: 22
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Best Local Similarity
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Best Local Similarity
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                                                                                                               APPLICANT: FEARON, Karen L.
APPLICANT: DINA, DINA,
APPLICANT: DINA, DINA
TITLE OF INVENTION: IMMUNOMODULATORY POLYNUCLEOTIDES AND
TITLE OF INVENTION: METHODS OF USING THE SAME
FILE REFERENCE: 377882001800
CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
FRIOR APPLICATION NUMBER: 60/258,675
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR APPLICATION NUMBER: 2000-12-27
PRIOR FILING DATE: 2000-12-27
                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR FILING DATE: 2000-12-27
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SOFTWARE: FastSEQ for Windows Version 4.0
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NUMBER OF SEQ ID NOS: 133
                                                                                                           NUMBER OF SEQ ID NOS: 133
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FEATURE:
                 ORGANISM: Artificial Sequence
                                    TYPE: DNA
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100.0%; Prr
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; OTHER INFORMATION: n = 5-bromocytosine US-10-033-243-46
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SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 46
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                   NUMBER OF SEQ ID NOS: 133
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 59
LENGTH: 22
                                                                                                                                                                                                                                                                                 Sequence 59, Application US/10033243
Publication No. US20030049266A1
GENERAL INFORMATION:
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Publication No. US20030049266A1
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                            TITLE OF INVENTION: IMMUNOMODULATORY POLYNUCLEOTIDES AND TITLE OF INVENTION: METHODS OF USING THE SAME FILE REFERENCE: 377882001800
CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR FILING DATE: 2000-12-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/033,243
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/258,675
PRIOR FILING DATE: 2000-12-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: IMMUNOMODULATORY POLYNUCLEOTIDES AND TITLE OF INVENTION: METHODS OF USING THE SAME FILE REFERENCE: 377882001800
                                                                                                                                                                                                                                              APPLICANT: FEARON, Karen L. APPLICANT: DINA, Dino
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FANTURE:
LOCATION: 5
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ORGANISM: Artificial Sequence
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LOCATION: 2, 5
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 Sequence 18, Appl Sequence 12, Appl Sequence 12, Appl Sequence 12, Appl Sequence 2448, Appl Sequence 2448, Appl Sequence 2455, App Sequence 563, Appl Sequence 576, Appl Sequence 576, Appl Sequence 576, Appl Sequence 577, Appl Sequence 11, Appl Sequence 637, Appl Sequence 637, Appl Sequence 11, Appl Sequence 125, Appl Sequence 1348, Appl Sequence 149, Appl Sequence 153, Appl Sequence 153, Appl Sequence 17, Appl Sequence 17, Appl Sequence 18, Appl Sequence 17, Appl Sequence 17, Appl Sequence 18, Appl Sequence 17, Appl Sequence 18, Appl Sequence 17, Appl Sequence 18, Appl Sequence 18, Appl Sequence 19, Appl Sequence 27, Appl Sequ
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JS-09-235-742-19

Sequence 19, Appli Patent No. 6498148

Application US/09235742

ALIGNMENTS

GENERAL INFORMATION:

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APPLICANT: Raz, Eyal
TITLE OF INVENTION: Immunization-Free Methods for Treating
TITLE OF INVENTION: Antigen-Stimulated Inflammation in a Mammalian Hos
TITLE OF INVENTION: Shifting the Host's Antigen Immune Responsiveness
TITLE OF INVENTION: Phenotype
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CURRENT FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR FILING DATE: 2000-03-28
PRIOR FILING DATE: 2000-05-16
PRIOR FILING DATE: 2000-05-16
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/820,484
CURRENT FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SOPTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 1
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  Query Match
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Patent No. 653406
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APPLICANT:
APPLICANT:
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APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Method for Increasing a Cytotoxic
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: 06510-188US1
                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Method for Increasing a Cytotoxic TITLE OF INVENTION: Lymphocyte Response in vivo. FILE REFERENCE: 06510-188US1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Raz, Eyal
                                                             OTHER INFORMATION: phosphorothioate ISS-ODN
                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
OTHER INFORMATION: Disulfide-linked phosphorothioate ISS-ODN
NAME/KEY: modified base
LOCATION: (1)...(1)
OTHER INFORMATION: disulfide thymine
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                     FEATURE:
                                                                                                                                               LENGTH: 22
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5, 6534062
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Horner, Anthony A.
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  Score 15;
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Length 22;
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GAACGTTCGAGATGA 20

GAACGTTCGAGATGA 22

Query Match Best Local Similarity Watches 15; Conserva

Conservative

71.4%; Score 15; 100.0%; Pred. No. cive 0; Mismatc

DB 4;

Length 22; Indels

Mismatches

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OTHER INFORMATION: Recombinant or Synthetic Sequence US-09-235-742-19

NUMBER OF SEQ ID NOS: 20 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 19 LENGTH: 22

TYPE: DNA
ORGANISM: Artificial Sequence

FEATURE:

FILE REFERENCE: 6510-170CON4
CURRENT APPLICATION NUMBER: US/09/235,742
CURRENT FILING DATE: 199-01-21
EARLIER APPLICATION NUMBER: 08/927,120
EARLIER FILING DATE: 1997-09-05
EARLIER APPLICATION NUMBER: 08/53,554
EARLIER FILING DATE: 1996-01-30
EARLIER APPLICATION NUMBER: 08/725,968
EARLIER FILING DATE: 1996-10-04
EARLIER APPLICATION NUMBER: 60/028,118
EARLIER FILING DATE: 1996-10-11

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FILE REFERENCE: 30448.64US01
CURRENT APPLICATION NUMBER: US/09/347,343A
CURRENT FILING DATE: 1999-07-02
NUMBER OF SEQ ID NOS: 40
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3:
SEQ ID NO 3:
LENGTH: 22
TYPE: DNA
ORGANISM: synthetic oligonucleotide
US-09-347-343-32
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Patent No. 6514948
GENERAL INFORMATION:
APPLICANT: RAZ, EYAL R.
APPLICANT: KOBAYASHI, Hiroko
TITLE OF INVENTION: METHOD FOR ENHANCING AN IMMUNE RESPONSE
                                                                     Query Match
Best Local Similarity
Matches 15; Conser
                                  GAACGTTCGAGATGA 20
                                                                       Conservative
                                                                                      71.4%; Score 15; 100.0%; Pred. No.
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US-08-776-251-10
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US-09-774-403A-1
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Best Local Similarity
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APPLICANT:
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CURRENT FILING DATE: 2002-04-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Tomoko Hayashi
APPLICANT: Dennis Camsoon
TITLE OF INVENTION: Immunomodulatory Polynucleotides in
TITLE OF INVENTION: Treatment of Infection by an Intracellular Pathogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Springer, Carol.
APPLICANT: Marais, Richard
TITLE OF INVENTION: Surfac
                                                             FILING DATE: 31-JAN-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB95/01782
                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/776,251
                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                             CITY: Arlington
STATE: Virginia
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                                                                                                                                                                                                                               COUNTRY:
APPLICATION NUMBER: GB 9415167.7 FILING DATE: 27-JUL-1994
                                                   FILING DATE:
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Antonio Catanzaro
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00 No. 6025340th Glebe Road, 8th Floor
                                                 27-JUL-1995
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US-08-776-251-10
                                                                            Query Match
Best Local :
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Best Local Similarity
                                                              Matches
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APPLICATION NUMBER: PCT/GB95/01782

PILING DATE: 27-UTL-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: GB 9415167.7

PILING DATE: 27-UTL-1994

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 25
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                 REGISTRATION NUMBER: 25
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 31-JAN-1997
                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 816 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                      TOPOLOGY: 1:
MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 816 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                           Local Similarity
                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                            NAME: Arthur R. Crawford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Arlington
STATE: Virginia
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                      TCGAACGTTCGAGA 17
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TCGAACGTTCGAGA 788
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1100 No. 6025340th Glebe Road, 8th Floor
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100.0%;
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Pred. No.
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US-09-060-756-260
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                                                                                        RESULT 10
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APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
                                Sequence 260, Application US/09060756 Patent No. 6183957
                  GENERAL
                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09103840A Patent No. 6294328
                                                                                                                                                                                                                    Matches
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Best Local
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Best Local Similarity
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APPLICANT: Cole, Stewart
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DIA SEQUENCES FOR STRAIN ANALYSIS IN
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT FILING DATE: 1998-06-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: FLEISCHMAN, Robert APPLICANT: WHITE, Owen R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: FLEISCHMAN, Robert D. APPLICANT: WHITE, Owen R.
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OTHER INFORMATION: "n" bases at various
OTHER INFORMATION: represent a, t, c or
                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
OTHER INFORMATION: H37Rv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                ENGTH: 4411529
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                INFORMATION:
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                                                                                                                                             TCGTCGAACGTTCG 733602
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                                                                                                                                                                                                                                                    Length 4411529;
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GENERAL INFORMATION:

APPLICANT: Cole, Stewart
APPLICANT: Buchrisser-Brosch,
APPLICANT: Gordon, Stephen
APPLICANT: Billault, Alain

Sequence 260, Application US/09670314 Patent No. 6492506

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RESULT 12
US-09-670-314-260
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; OTHER INFORMATION: anti-Rh(D)
US-09-240-274-197
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US-09-240-274-197/c
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SEQ ID NO 197
LENGTH: 321
TYPE: DNA
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SEQ ID NO 260
LENGTH: 321
TYPE: DNA
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                                                                                                                                                                     Query Match
Best Local Similarity
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Matches 1
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TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT FILING DATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
                                                                                                                                                                                                                                                                                                                                                                          EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
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APPLICANT: Gordon, Stephen
APPLICANT: Billault, Alain
TITLE OF INVENTION: METHOD FOR ISOLATING A POLYNUCLEOTIDE OF INTEREST FROM
TITLE OF INVENTION: THE GENOME OF A MYCOBACTERIUM USING A BAC-BASED DNA
TITLE OF INVENTION: LIERARY APPLICATION TO THE DETECTION OF MYCOBACTERIA
FILE REFERENCE: 3495-0169
CURRENT APPLICATION NUMBER: US/09/060,756
CURRENT FILING DATE: 1998-04-16
NUMBER OF SEQ ID NOS: 743
                                                                                                                                                                                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: 60/028,550
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NAME/KEY: unsure
LOCATION: (vario)
                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
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                                                                           292 CGAACGTTCGAGA 280
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                                                                                                                                                      13;
                                                                                                5 CGAACGTTCGAGA 17
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                                                                                                                                                                     61.9%;
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Pred. No.
                                                                                                                                                                                                                                              antibody clone SH8
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US-09-252-991A-16046
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ORGANISM: Mycobacterium tuberculosis
FEATURE:
NAME/KEY: unsure
LOCATION: (various positions within
OTHER INFORMATION: applicants are un
US-09-670-314-260
                                                                                                                                                                                                                                 US-09-252-991A-9162/c
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SEQ ID NO 16046
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                                                                                                                                                                                      Sequence 9162, Application US/09252991A Patent No. 6551795
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                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local
APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD FOR ISOLATING A POLYNUCLECTIDE OF INTEREST FROM TITLE OF INVENTION: THE GENOME OF A MYCOBACTERIUM USING A BAC-BASED DNA TITLE OF INVENTION: LIBRARY APPLICATION TO THE DETECTION OF MYCOBACTERIA FILE REFERENCE; 3495-0169
CURRENT APPLICATION NUMBER: US/09/670,314
CURRENT FILING DATE: 2001-01-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/060,756
PRIOR FILING DATE: 1998-04-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 462
TYPE: DNA
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100.0%; Pred. No.
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27;
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FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 16554
LENGTH: 762
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US-09-252-991A-16554/c
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US-09-252-991A-9246/c
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                                                                                                                                                                       GEMERAL INFORMATION:
GEMERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
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SEQ ID NO 9246
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NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 9162
LENGTH: 573
                                                                                                                                                                                                                                                                                                               Sequence 16554, Application US/09252991A Patent No. 6551795
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 663
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
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NAME/KEY: unsure
LOCATION: (17)
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ORGANISM: Pseudomonas aeruginosa
ORGANISM: Pseudomonas aeruginosa
                      DNA
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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Pred. No.
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26;
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US-09-107-532A-1566
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                                                                                                                            SEQUENCE DESCRIPTION: SEQ.ID NO: US-09-107-532A-1566
                                                             Query Match
Best Local S
Matches 13
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GENERAL INFORMATION:
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Patent No. 6583275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051871
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                        TOPOLOGY: circular MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                          ANTI-SENSE: NO
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Similarity 100.0%; P
13; Conservative 0;
GTCGAACGTTCGA 253
                              GTCGAACGTTCGA 15
                                                                                                                                                                                                                                                                                                         LENGTH: 813 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                                                                             ORGANISM: Enterococcus faecium
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                                                              Conservative
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(B) LOCATION 1...813
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AND AMINO ACID SEQUENCES RELATING TO
FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
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US-09-252-991A-9074/c
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US-09-252-991A-8921
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APPLICANT: MARC J. Rubenfield et al.
APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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NUMBER OF SEQ ID NOS:
SEQ ID NO 8921
LENGTH: 1221
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PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 9074
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APPLICANT: Marc J.
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                                                                  Query Match
Best Local Similarity
Matches 13; Conserv
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
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TITLE OF INVENTION:
TITLE OF INVENTION:
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LOCATION: (458)
OTHER INFORMATION:
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ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                        NAME/KEY: unsure
LOCATION: (951), (1078)
OTHER INFORMATION: Ide
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212
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                        6 GAACGTTCGAGAT 18
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GAACGTTCGAGAT 200
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1998-02-18
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
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RESULT 20 US-09-252-991A-8710 ; Sequence 8710, Application US/09252991A

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RESULT 22
US-09-252-991A-15954
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GENERAL INFORMATION:
APPLICANT: MARC J. Rubenfield et al.
APPLICANT: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 16328
LENGTH: 1986
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                     Sequence 15954, Application US/09252991A Patent No. 6551795
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LENGTH: 1545
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATITIES OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS A
FILE REFERENCE: 107196.136
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 1998-07-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 13; Conserv
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                                                                                                                                                                                                                                                                   1676
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                                                                                                                                                                                                                                                                                                      ACGTTCGAGATGA 20
                                                                                                                                                                                                                                                                 ACGTTCGAGATGA 1664
                                                                                                                                                                                                                                                                                                                                                 61.9%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61.9%; Score 13; ilarity 100.0%; Pred. No. Conservative 0; Mismatci
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US 60/094,190
                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                         DB 4;
24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D SEQUENCES RELATING TO PSEUDOMONAS AND THERAPEUTICS
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GENERAL INFORMATION:
APPLICANT: Baker, Adam
APPLICANT: Chicca, Matthew
APPLICANT: Chicca, Susanna
APPLICANT: Kurbauer, Robert
APPLICANT: Schaffner, Gotthold
TITLE OF INVENTION: Chicken Embryo Lethal Orphan (CELO) Virus
FILE REFERENCE: 0652.1800000
CURRENT APPLICATION NUMBER: US/09/171,461
CURRENT FILING DATE: 1999-01-12
EARLIER APPLICATION NUMBER: PCT/EP97/01944
EARLIER FILING DATE: 1997-04-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ:ID NO 19954
ENGTH: 2091
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US-09-171-461-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 1
LENGTH: 43804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1,
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                                                                                                                                                                                                                                                         FEATURE:

NAME/KEY: polyA site

TOCATION: (17526)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
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                                                LOCATION: (21102)
OTHER INFORMATION: /gene:
                                                                                                                                                                                           NAME/KEY: gene
LOCATION: (17559)..(21754)
OTHER_INFORMATION: /gene: L3
                                                                                                                                                                                                                                                                                                                       NAME/KEY: gene
LOCATION: (15110)..(17495)
OTHER INFORMATION: /gene:
NAME/KEY: misc_feature
LOCATION: (21123)
                                                                     NAME/KEY: misc_feature
LOCATION: (21102)
                                                                                                                           OTHER INFORMATION:
                                                                                                                                                         FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: /note= L2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: gene
LOCATION: (121)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: CELO Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                          OCATION:
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les 13; Conserv
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                                                                                                                        /gene:
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                                                    L3
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                                                    /note= protease
                                                                                                                           /note=
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                                                                                                                         hexon splice acceptor site
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                                                    splice acceptor site
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OTHER INFORMATION: /gene:

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/note= protease splice acceptor site

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RESULT 24
US-09-286-098-11/c
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Patent No. 6218371
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
           APPLICANT: Krieg, Arthur M.
APPLICANT: Weiner, George
TITLE OF INVENTION: Methods and Products for Stimulating the
TITLE OF INVENTION: Immune System Using Immunotherapeutic Oligonucleotides
TITLE OF INVENTION: Cytokines
TITLE REFERENCE: C1039/7026/HCL
CURRENT APPLICATION NUMBER: US/09/286,098
CURRENT FILING DATE: 1999-04-02
EARLIER APPLICATION NUMBER: US 60/080,729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: gene
LOCATION: (23680)..(27886)
OTHER INFORMATION: /gene: L4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: polyA_site
LOCATION: (21824)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: polyA_site
LOCATION: (31770)
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LOCATION: (28363)..(31768)
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LOCATION: (28341)
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JOCATION: (28315)
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OCATION: (21836)
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                                                                                                                                                                                                                                                                                                             25862
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FILING DATE:
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                                                                                                                                                                                                                                                                                                             GAACGITCGAGAT 25874
                                                                                                                                                                                                                                                                                                                                               GAACGTTCGAGAT 18
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                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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1998-04-03
                                                                                                                                                                                                                                                                                                                                                                                                    61.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                    score 13;
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                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
20;
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RESULT 25
US-09-325-193A-91/c
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US-08-882-704A-18
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LENGTH: 20
TYPE: DNA
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 6406705
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                 Patent No. 5879906
GENERAL INFORMATION:
                                                                                                                                                                                               Sequence 18, Application US/08882704A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Schorr, Joachim
APPLICANT: Krieg, Arthur M.
TITLE OF INVENTION: Use of Nucleic Acids Containing
TITLE OF INVENTION: Unmethylated CpG Dinucleotide as
FILE REFERENCE: C1039/7025/HCL
CURRENT APPLICATION NUMBER: US/09/325,193A
CURRENT FILING DATE: 1999-06-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1998-
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1997-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 1999-06-03
PRIOR APPLICATION NUMBER: US 09/154,614
PRIOR FILING DATE: 1998-09-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Synthetic Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20
                                                                     APPLICANT: Leader, Mich
TITLE OF INVENTION: GLI
NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                     APPLICANT: Jefferson, Richard A
APPLICANT: Wilson, Katherine J.
APPLICANT: Leader, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                            STREET:
COUNTRY:
                                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                        12;
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12; Conservative
                                                                                                                                                                                                                                                                                                        6 GAACGTTCGAGA 17
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           Seattle
Washington
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                                             6300
                                                                                                                                                                                                                                                                                                                                        Conservative
USA
                                            SEED and BERRY LLP 00 Columbia Center,
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                                                                                                                                                     Richard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT/US98/04703
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Pred. No.
                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                        REPRESSORS
                                             701
                                               Fifth Avenue
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hes 0;
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                                                                                                          AND
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Best Local Similarity
Matches 12; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-882-704A-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18, Application US/08882704A Patent No. 5879906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5879906
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (206) 682-6031 INFORMATION FOR SEQ ID NO:
                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6931
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/882,704A
FILING DATE: 25-JUN-1997
CLASSIFICATION: 435
ATTORNEY JACTEME 135
ATTORNEY JACTEME 135
                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FORM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION WIMBER: US/08/882,704A
FILING DATE: 25-UN-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: No. 5879906tenburg Ph.D., Carol
REGISTRATION NUMBER: 39,317
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: NO. 5879906tenburg Ph.D., Carol
REGISTRATION NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 190106.404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jefferson, Richard A. APPLICANT: Wilson, Katherine J. APPLICANT: Leader, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TLE OF INVENTION:
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 98104-7092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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  STRANDEDNESS:
                                           ENGTH:
nucleic acid
EDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           57.1%; Score 12; DB 2; llarity 100.0%; Pred. No. 1.3e+(Conservative 0; Mismatches
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US-09-151-957-18/c
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US-09-151-957-18
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                                                  Sequence 18, Application US/09151957
Patent No. 6429292
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18, Application US/09151957
Patent No. 6429292
GENERAL INFORMATION:
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Best Local :
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TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PROJECTION NUMBER: US/09/151,957
FILING DATE: 11-Sep-1998
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/882,704
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: No. 6429292tenburg Ph.D., Carol
REGISTRATION NUMBER: 39,317
REGISTRATION NUMBER: 39,317
                                                                                                                                                                                                                                                      Local Similarity 100.0%; I les 12; Conservative 0;
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:
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ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center,
CITY: Seattle
APPLICANT: Jefferson, Richard A.
Wilson, Katherine J.
Leader, Michael
                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 18:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
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                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 22 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
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Leader, Michael
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                                                                                                                                                                                                                                                      57.1%; Score 12; DB 100.0%; Pred. No. 1.: tive 0; Mismatches
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                                                                                                                                                                                                                                                                                          DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                                                                                                                                                    APPLICANT: KRAGH, KARSTEN
APPLICANT: BOJKO, MAJA
APPLICANT: NIELSEN, JOHN
APPLICANT: MARCUSSEN, JAN
TITLE OF INVENTION: ALPHA-1,4-GLUCAN LYASE FROM
TITLE OF INVENTION: A FUNGUS INFECTED ALGAE, ITS PURIFICATION
NIMBER OF CECUTENCES: A FUNGUS INFECTED ALGAE, ITS PURIFICATION
                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                       CITY: Newport Beach
STATE: CA
COUNTRY: U.S.A.
                                    OPERATING SYSTEM:
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                                                                                                                                                                                                                              ADDRESSEE:
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FILING DATE: 11-Sep-1998
CLASSIEICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/882,704
FILING DATE: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: No. 642922tenburg ph.D., Carol
REGISTRATION NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 190106.404
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 682-6931
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
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ADDRESSEE: SEED and BERRY LLP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
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NRE: FastSEQ Version 1.5
APPLICATION DATA:
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L2; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 22 base pairs
                                                                                                                                                                                                        E: Knobbe, Martens, Olson & Bear
620 Newport Center Drive 16th Floor
                                                                                                                                                                                                                                                                                                                                                                                                               BOJSEN, KIRSTEN
                                                          IBM Compatible
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RESULT 31
US-08-399-412A-58/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                       OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/399,412A
FILING DATE: 6-MARCH-1995
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
TYPE: nucleic acid
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FILING DATE: 15-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                        APPLICATION NUMBER: 07/53
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: High-Affinity Oligonucleotide
TITLE OF INVENTION: Ligands To Secretory Phospholipase
TITLE OF INVENTION: A2 (spla2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: Genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 714-760-0404
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                                                    FILING DATE: 21-OCTOBER-: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Altman, Daniel E
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: DY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Englewood
STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 02-JUI
                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 80111
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TELEFAX: 714-760-9502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parma, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                         IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Swanson & Bratschun, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                          Diskette,
                              L. Bernard
                                                                     21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-1996
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NUMBER:
                                                                                        07/964,624
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                                                                                                                                             07/536,428
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                                                                                                                                                                                                                                                                                                                                                                                          3.5 inch, 1.44 MB storage
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TELECOMMUNICATION INFORMATION:

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Query Match
Best Local Similarity
Thes 12; Conserve
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; Sequence 14, Application US/08633768A
; Patent No. 6013504
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                                                                            US-08-633-768A-14
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 77 base pairs
                                                                                                                                                                  NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 9321301.5
FILING DATE: 15-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 02-JUL-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ALPHA-1,4-GLUCAN LYASE FROM
TITLE OF INVENTION: A FUNGUS INFECTED ALGAE, ITS PURIFICATION
NUMBER OF SEQUENCES: 25
                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: NIELSEN, JOHN APPLICANT: MARCUSSEN, JAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT:
                                                                                                                                                                                                                                                           NAME: Altman, Daniel E
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: DY
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FABLSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 111
                                                                                                                  STRANDEDNESS: single
                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                               TELEPHONE: 714-760-04
TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
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Local Similarity 100.0%;
es 12; Conservative (
                                                                                                       TOPOLOGY:
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Newport Beach
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BOJSEN, KIRSTEN
KRAGH, KARSTEN
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O Newport Center Drive 16th Floor
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                                                                                          Genomic DNA
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                            57.1%; Score 12;
100.0%; Pred. No.
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; Pred. No.
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                              1.1e+02;
                                           DB 3;
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nes 0; Indels
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RESULT 35
US-09-313-294A-2857
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; OTHER INFORMATION: Incyte ID No. 6476212 700552684H1
US-09-313-294A-2448
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US-09-313-294A-2448
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; ORGANISM: Acinetobacter baumannii
US-09-328-352-3855
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US-09-328-352-3855/c
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SEQ ID NO 2448
LENGTH: 212
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SEQ ID NO 3855
LENGTH: 186
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APPLICANT: Gary L.
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                                                                                                                                    Best Local Similarity Matches 12; Conserv
                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN
FILE REFERENCE: PL-0017 US
                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/313,294A CURRENT FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lalgudi, Raghunath V. APPLICANT: Ito, Laura Y.
                                                                                                                                                                                                                                                                         ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                         FEATURE:
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nes 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09313294A
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                                                                                                                                      Conservative
                                                                                                                                                  57.1%;
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                                                                                                                                                    Score 12;
Pred. No.
                                                                                                                                                                  DB 4; Length 212;
                                                                                                                                                    1.1e+02;
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Sequence 2857, Application US/09313294A Patent No. 6476212 GENERAL INFORMATION:

APPLICANT:

Ito, Laura Y. Sherman, Bradley K.

Lalgudi, Raghunath V

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; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte II
; NAME/KEY: unsure
; LOCATION: 257
; OTHER INFORMATION: a, t, c,
US-09-313-294A-2857
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                                                                                                                                                                                    RESULT 37
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APPLICANT: Ito, Laura Y.

APPLICANT: Sherman, Bradley K.

TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR
FILE REFERENCE: PL-0017 US

CURRENT APPLICATION NUMBER: US/09/313,294A

CURRENT FILING DATE: 1999-05-14

NUMBER OF SEQ ID NOS: 7600

SOFTWARE: PERL Program

SEQ ID NO 4896

LENCTH: 283
                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 12
                                                                                      Sequence 69, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
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Best Local
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SEQ ID NO 2857
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LOCATION: 122, 140
OTHER INFORMATION: a, t,
-09-313-294A-4896
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             APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
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OTHER INFORMATION: Incyte ID No.
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12; Conserv
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Conservative
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US-09-252-991A-5259/c
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; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-69
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PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 69
TLENGTH: 288
                                                      SOFTWARE: PatentIn Ver.
SEQ ID NO 563
LENGTH: 406
                                                                                                                 GENERAL INFORMATION:

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Cole, Stewart

APPLICANT: Buchrieser-Brosch, Roland

APPLICANT: Buchrieser-Brosch, Roland

APPLICANT: Billault, Alain

ITILE OF INVENTION: METHOD FOR ISOLATING A POLYNUCLEOTIDE OF INTEREST FROM

TITLE OF INVENTION: THE GENOME OF A MYCOBACTERIUM USING A BAC-BASED DNA

TITLE OF INVENTION: LIBRARY APPLICATION TO THE DETECTION OF MYCOBACTERIA

FILE REFERENCE: 3495-0169

CURRENT APPLICATION UMDBER: US/09/080,756

CURRENT APPLICATION UMDBER: US/09/080,756

CURRENT APPLICATION UMDBER: US/09/080,756

CURRENT FILING DATE: 1998-04-16

NUMBER OF SEQ ID NOS: 743
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SEQ ID NO 5259
LENGTH: 378
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Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18
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ORGANISM: Pseudomonas aeruginosa
FEATURE:
                 TYPE: DNA ORGANISM: Mycobacterium tuberculosis
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           NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 3530
LENGTH: 432
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
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Best Local Similarity
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-09-252-991A-3530
                                                                                                                      FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR TILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                                                                                                   APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
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CURRENT FILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 09/060,756
PRIOR FILING DATE: 1998-04-16
NUMBER OF SEQ ID NOS: 743
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: METHOD FOR ISOLATING A POLYNUCLEOTIDE OF INTEREST FROM TITLE OF INVENTION: THE GENOME OF A MYCOBACTERIUM USING A BAC-BASED DNA TITLE OF INVENTION: LIBRARY APPLICATION TO THE DETECTION OF MYCOBACTERIA PILE REFERENCE: 3495-0169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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APPLICANT: Buchr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: unsure LOCATION: (various positions within the sequence) OTHER INFORMATION: applicants are uncertain of bases designated as "n"
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Local Similarity 100.0%; Pred. No. 1.:
1es 12; Conservative 0; Mismatches
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100.0%; Pred. No. 1.1e+02;
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hes 0;
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 69
LEGATH: 658
TYPE: DNA
ORGANISM: Usnea florida
US-08-861-774E-69
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US-08-861-774E-69/c
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US-09-252-991A-4565
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Best Local Similarity
Matches 12; Conserv
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APPLICANT: Waters, Barbara
APPLICANT: Miao, Vivian
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SEQ ID NO 4565
LENGTH: 441
                                                                  Matches
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Best Local :
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                                                                                   Query Match
Best Local
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CURRENT FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 94
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TITLE OF INVENTION: NUCLEIC ACID AND ANINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR PRIOR DATE: 1998-07-27

NUMBER OF SEC ID NOC. 23142
                                                                                                                                                                                                                                                                                                  APPLICANT: TONG, SECW TITLE OF INVENTION: METHOD FOR ISOLATION OF BIOSYNTHESIS TITLE OF INVENTION: BIOACTIVE MOLECULES FILE REFERENCE: 9993-006
                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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ORGANISM: Pseudomonas aeruginosa
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                                                                                     Local Similarity
187
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                               10 GTTCGAGATGAT 21
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GTTCGAGATGAT 176
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                                                                  ilarity 100.0%;
Conservative 0;
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100.0%; Pred. No. 1.1e+02;
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0; Mismatches
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Sequence 78,

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RESULT 45
US-09-107-532A-1566/c
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; OTHER INFORMATION: brCBF3 gene
US-09-198-119C-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
SOFTWARE: PATENTIN VE:
SEQ ID NO 78
LENGTH: 806
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1566, Application US/09107532A
PATCHI NO. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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PRIOR APPLICATION NUMBER: US 09/018,234
PRIOR FILING DATE: 1998-02-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1998-02-03
PRIOR APPLICATION NUMBER: US 09/017,575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Brassica rapa
                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                            ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 7310 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     692 ACGTTCGAGATG 703
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APPLICATION NUMBER: 60/085,598
                                                                                                              OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
                                                                                                                                                           COMPUTER: PC
                                                                                                                                                                                                                                                                            STATE: Massachusetts
                                                                                                                                                                                                                                                                                                 STREET: 100 Beaver Street CITY: Waltham
                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stockinger, Eric
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Jiang, Cai-Zhong
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AND AMINO ACID SEQUENCES RELATING TO
FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 806;
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US-09-198-119C-84
                                                                                                                                             PRIOR APPLICATION NUMBER: 1
PRIOR FILING DATE: 1998-02:
PRIOR APPLICATION NUMBER: 1
PRIOR FILING DATE: 1998-02:
NUMBER OF SEQ ID NOS: 95
                                                                                                     SOFTWARE: 1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Plant Having Altered Environmental Stress Tolerance FILE REFERENCE: 19117.713 Seq List CURRENT APPLICATION NUMBER: US/09/198,119C CURRENT FILING DATE: 1998-11-23 CURRENT FILING DATE: 1998-11-23 PRIOR APPLICATION NUMBER: US 08/706,270
                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 08/706,270
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: US 09/018,233
PRIOR FILING DATE: 1998-02-03
PRIOR APPLICATION NUMBER: US 09/017,816
PRIOR APPLICATION NUMBER: US 09/018,235
PRIOR APPLICATION NUMBER: US 09/018,235
PRIOR APPLICATION NUMBER: US 09/018,235
                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1998-PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-
                                      LENGTH: 830
TYPE: DNA
ORGANISM: Brassica rapa
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OTHER INFORMATION: brCBF6 gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     253 TCGAACGTTCGA 242
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Jaglo-Ottosen, Kirsten
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Jiang, Cai-Zhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/09198119C
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                                                                                                                                                                                                            IUMBER: US 09/018,227
1998-02-03
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IUMBER: US 09/017,575
1998-02-03
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                                                                                                                                                                                              US 09/018,234
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US-09-252-991A-5231/c
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                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
SEQ ID NO 5231
LENGTH: 981
                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION: APPLICANT: Marc J.
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Patent No. 6551795
GENERAL INFORMATION:
                                                                                          Query Match
Best Local (
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: MARC J. RUBENfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/094,190 PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                  TYPE: DNA ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Pseudomonas aeruginosa
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22
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                                                                                          Similarity
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GTCGAACGTTCG 11
                                    GTCGAACGTTCG 14
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                                                                          Conservative
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                                                                                          57.1%; Score 12; 100.0%; Pred. No.
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1e+02;
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le+02;
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TITLE OF INVENTION: MUCLEIC ACID AND AMINO A TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTI FILE REFERENCE: 107195.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 3539
LENGTH: 1002
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
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US-09-091-097-37/c
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Sequence 37, App
Sequence 37, App
64324
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                                                   PILING DATE:
CLASSIFICATION: 424
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WEINER, MARC S.
REGISTRATION NUMBER: 32,181
REFERENCE/DOCKET NUMBER: 1427
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTIGENIC PROTEIN ORGINATING TITLE OF INVENTION: MALASSEZIA NUMBER OF SEQUENCES: 58 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Marc J. Rubenfield et al. 
VENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS 
VENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                        E: BIRCH, STEWART, KOLASCH & BIRCH, LLP
PO BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AKIYAMA, KAZUO
YASUEDA, HIROSHI
YAMAGUCHI, HIDEYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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ONISHI, YOSHIMI
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                                                                                                                                                                                                       us/09/091,097
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                                                                                             1422-0346P
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                                                                                                                                                                                                                                           Version #1.30
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1e+02;
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TYPE: nucleic acid

STRANDEDNESS: double

TOPOLOGY: linear

MOLECULE TYPE: CDNA to mRNA
US-09-091-097-37
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                                                                       US-09-252-991A-16487
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US-09-252-991A-16116
                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                       SEQ ID NO 16487
LENGTH: 1038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 16116
LENGTH: 1038
Best Local Similarity Matches 12; Conserv
                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                          Sequence 16487, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Patent No. 655179
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PRIOR APPLICATION NUMBER: US 6
PRIOR APPLICATION DATE: 1998-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR TILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/252,991A.
CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1998-07-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                  TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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 Conservative
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                                                                                                                                                                                                            TUMBER: US 60/074,788
1998-02-18
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                57.1%; Score 12; 100.0%; Pred. No.
                                                                                                                                                               33142
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 Mismatches
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                                 DB 4;
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1e+02;
                    1e+02;
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                                  Length 1038;
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US-09-252-991A-6391/c
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US-08-229-287-1
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US-08-229-287-1/c
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Sequence 6391, Application US/09252991A
Patent NO. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: AERUGINOSA FOR DIJ
TITLE OF INVENTION: AERUGINOSA FOR DIJ
                                                                                                                                                                                                                                  Matches
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CLASSIFICATION: 435
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 08/
APPLICATION DATA:
FILING DATE: US 07/
APPLICATION UNMBER: US 07/
FILING DATE: 08-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415/857-1125
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1113 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/229,287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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APPLICANT: Murry, Lynn B.
APPLICANT: Scarafii
                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 415/354-3588
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Marcus-Wyner, Lynn
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                NAME/KEY:
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                                                                                                                                                                   695 GTTCGAGATGAT 684
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                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
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975 California Avenue
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VENTION: VIRUS RESISTANT CORN PLANTS
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                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                              single
 Rubenfield et al.

NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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; Pred. No.
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                                                                                                                                                                                                                                   Mismatches
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. le+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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CURRENT APPLICATION NUMBER: 105/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
                                                                                                                                             TELEFAX: 302-773-0164
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: HITZ, WILLIAM DEAN
TITLE OF INVENTION: NUCLEOTIDE.
TITLE OF INVENTION: STERASE GE
TITLE OF INVENTION: THE REGULAT
TITLE OF INVENTION: CONTENT OF
TITLE OF INVENTION: AND CANOLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Pseudomonas aeruginosa
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ANTI-SENSE:
                              MOLECULE TYPE:
                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 1303 base pairs
TYPE: nucleic acid
                                                                                                                                                                             NAME: CHRISTENBURY, LYNNE M.
REGISTRATION NUMBER: 30,971
REFERENCE/DOCKET NUMBER: CR-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-992-5481
                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/10627
FILING DATE: AUGUST 25, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 3
                  HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: B. I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
CITY: WILMINGTON
STATE: DELAWARE
                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE:
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                                                        TOPOLOGY:
                                                                     STRANDEDNESS:
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5. 5955650
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                                                                       single
                                   DNA (genomic)
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100.0%; Pred. No.
tive 0; Mismatc
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MESULT 58
US-09-252-991A-6308/c
US-09-252-991A-6308/c
; Sequence 6308, Application US/09252991A
; Patent No. 6551795
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US-09-252-991A-64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; SEQ ID NO 943
; LENGTH: 1362
; TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 64, Applicat Patent No. 6551795 GENERAL INFORMATION:
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APPLICANT: Gary L. Breton et al.
APPLICANT: INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                    Matches
                                                                                                                                                                                     Best
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                   SEQ ID NO 64
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR RIPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
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CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
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                                                                                                                                                                                                                                             LENGTH: 1368
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09252991A
                                                                                                                                                                    Conservative
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Mismatches
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GENERAL

APPLICANT:

INFORMATION: NT: Marc J.

Rubenfield et al.

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RESULT 60
US-09-677-488A-7/c
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; LCCATION: (0)...(0)
; OTHER INFORMATION: permease, fully spliced cDNA
US-09-351-224E-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                 Sequence 7, Application US/09677488A Patent No. 6482621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 7
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APPLICANT: Maddox, Joyce
APPLICANT: Gilliam, Jacob
APPLICANT: Folkerts, Otto
APPLICANT: Crasts, Oswald R.
TITLE OF INVENTION: Compositions and Methods for Fumonisin
TITLE OF INVENTION: Detoxification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 5718-111
CURRENT APPLICATION NUMBER: US/09/351,224E
CURRENT FILING DATE: 1999-07-12
NUMBER OF SEQ ID NOS: 11
                                                                          APPLICANT: Duvick, Jon
APPLICANT: Maddox, Jo;
APPLICANT: Gilliam, J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Crasta, Oswald R.
TITLE OF INVENTION: Compositions and Methods for Fumonisin
TITLE OF INVENTION: Detoxification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 1578
TYPE: DNA
ORGANISM: Exophiala spinifera
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No. 638817
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Gilliam, Jacob
Folkerts, Otto
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APPLICANT: Duvick, Jon
APPLICANT: Maddox, Joyce
APPLICANT: Maddox, Joyce
APPLICANT: Maddox, Jacob
APPLICANT: Folkerts, Otto
APPLICANT: Folkerts, Otto
APPLICANT: Crasta, Oswald R.
TITLE OF INVENTION: Compositions and Methods for Fumonisin
TITLE OF INVENTION: Detoxification
FILE REFERENCE: 35718/204101
CURRENT APPLICATION NUMBER: US/09/677,682B
CURRENT FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: 09/351,224
PRIOR FILING DATE: 1999-07-12
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Best Local Similarity
                                                                                                                  RESULT 62
US-09-252-991A-3548/c
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            Sequence 3548, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: AERUGINOSA FOR DIF
                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 1578
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CURRENT APPLICATION NUMBER: US/09/677,488A
CURRENT FILING DATE: 2000-10-02
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FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: permease, fully spliced cDNA
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NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: permease, fully spliced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 1578
TYPE: DNA
ORGANISM: Exophiala spinifera
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ORGANISM: Exophiala spinifera
FEATURE:
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          Rubenfield et al.

NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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Pred. No.
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GENERAL INFORMATION:
APPLICANT: Duvick, Jon
APPLICANT: Maddox, Joyce
APPLICANT: Gilliam, Jacob
APPLICANT: Folkerte, Otto
APPLICANT: Crasta, Oswald R.
TITLE OF INVENTION: Compositions and Methods for Fun
TITLE OF INVENTION: Detoxification
FILE REFERENCE: 35718/204100
CURRENT APPLICATION UMBER: US/09/677,488A
CURRENT FILING DATE: 2000-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Exophiala spinifera
FEATURE:
NAME/KEY: misc_feature
LCCATION: (0) ...(0)
OTHER INFORMATION: permease, partially spliced cDNA
US-09-351-224E-6
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US-09-351-224E-6/c
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                                                                                                                                                                                                                                                  RESULT 64
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CURRENT APPLICATION NUMBER: US/09/351,224E
CURRENT FILING DATE: 1999-07-12
NUMBER OF SEQ ID NOS: 11
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US OF PRIOR FILING DATE: 1998-07-27 NUMBER OF SEQ ID NOS: 33142 SEQ ID NO 3548
                                                                                                                                                                                           Sequence 6, Application US/09677488A Patent No. 6482621
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Best Local (
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Best Local Similarity
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APPLICANT:
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TITLE OF INVENTION: Compositions and Methods for Fumonisin
TITLE OF INVENTION: Detoxification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Maddox, Joyce
APPLICANT: Gilliam, Jacob
APPLICANT: Folkerts, Otto
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ORGANISM: Pseudomonas aeruginosa
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100.0%; Pred. No.
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Pred. No.
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RESULT 66
US-07-754-918A-1
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Best Local S
Matches 12
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SOFTWARE: FASTSEQ for
SEQ ID NO 6
                            Sequence 1, Application US/07754918A

Patent No. 5286484

GENERAL INFORMATION:
APPLICANT: Rodriguez,R.S. et al
TITLE OF INVENTION: NUCLEOTIDE SEQUENCE CODING FOR AN
TITLE OF INVENTION: OUTER MEMBRANE PROTEIN FROM NEISSERIA MENINGITIDIS AND USE
TITLE OF INVENTION: OF SAID PROTEIN IN VACCINE PREPARATIONS
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version SEQ ID NO 6
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APPLICANT: Maddox, Joyc
APPLICANT: Gilliam, Joc
APPLICANT: Folkerts, Oth
APPLICANT: Crasta, Oswa
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Best Local Similarity
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TITLE OF INVENTION: Compositions and Methods for Fumonisin TITLE OF INVENTION: Detoxification

FILE REFERENCE 35718/204101

CURRENT APPLICATION NUMBER: US/09/677,682B

CURRENT FILING DATE: 2000-10-02

PRIOR APPLICATION NUMBER: 09/351,224

PRIOR FILING DATE: 1999-07-12

NUMBER OF SEQ ID NOS: 11
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NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: permease, partially spliced cDNA
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TYPE: DNA
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wes 12; Conserv
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5. 6534291
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Folkerts, Otto
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Michaelson, Spivak and Wallace, Esq
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Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                      SEQ ID NO 1
LENGTH: 1900
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/09555000 Patent No. 6489108
                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/555,000 CURRENT FILING DATE: 2000-05-23 PRIOR APPLICATION NUMBER: PCT/US98/26971
                                                                                                                                                                                                                       PRIOR FILING DATE: 1:
                                                                                                                                                                                                                                                                                                            FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                            APPLICANT: Genencor International, Inc.
TITLE OF INVENTION: Proteases from Gram Positive Organisms
                                                                                                                                                                                                            SOFTWARE:
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                                                                                    NAME/KEY: CDS
LOCATION: (134)...(1774)
                                                                                                                        FEATURE:
                                                                                                                                        ORGANISM: Bacillus subtilis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: 9
ORIGINAL SOURCE:
ORGANISM: N. 1
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LENGTH: 1836 bp
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MEDIUM TYPE: 5 1/4" 360Kb IBM compatible diskette
COMPUTER: IBM ps/2 Model 80
OPERATING SYSTEM: MS-DOS 5.0
SOFTWARE: Microsoft Word 5.5
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)530-6671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Michaelson, Peter L.
REGISTRATION NUMBER: 30090
                                                                                                                                                       PE: DNA
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TYPE: NUCLEOTIDE WITH CORRESPONDING AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
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Road, P. O. Box 8489
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57.1%; Score 12;
100.0%; Pred. No.
tive 0; Mismatcl
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97;
                     DB 4;
97;
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US-08-358-117-1/c
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Patent No. 5608147
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Best Local Similarity
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                                                                                       GENERAL INFORMATION:
APPLICANT: STREBER, WOLFGANG R.
APPLICANT: TIMMIS, KENNETH N.
APPLICANT: ZENK, MEINHART H.
APPLICANT: ZENK, MEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION: NAME: Esmond, Robert W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kaphammer, Bryan J.
TITLE OF INVENTION: tfdA Gene S
TITLE OF INVENTION: Use Thereof
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                 CORRESPONDENCE ADDRESS
                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (202)
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FILING DATE: 15-DEC-1994
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1016 ACGTTCGAGATG 1005
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751..1611
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FROMMER LAWRENCE & HAUG LLP
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100.0%; Pred. No.
tive 0; Mismatc
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tfdA Gene Selectable Markers in Plants and
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

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                                                                                                                                              SOFTWARE: Pat
SEQ ID NO 3
LENGTH: 2426
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Matches
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                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atent No.
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 01
PRIOR FILING DATE: 1995-03-31
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                    TITLE OF INVENTION: GLUCAN HAVING CYCLIC STRUCTURE AND METHOD FOR PRODUCING THE SAN FILE REFERENCE: 9900-0002.20 CURRENT APPLICATION NUMBER: US/08/528,026C CURRENT FILING DATE: 1995-09-13
                                                                                                                                                                                                                                                                                                       APPLICANT: FUJII, I
                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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NFORMATION FOR SEQ ID NO: 1:
                                              NAME/KEY: misc feature
LOCATION: (385)..(392)
OTHER INFORMATION: S SD sequence
                                                                                                              TYPE: DNA
ORGANISM: Bacillus stearothermophilus (Strain TRBE14)
              NAME/KEY: misc_feature
LOCATION: (402)..(2357)
OTHER INFORMATION: P CDS
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LENGTH: 2058 base pair
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NAME: LAWRENCE, WILLIAM F.
REGISTRATION NUMBER: 28,029
REFERENCE/DOCKET NUMBER: 514413-3526
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-588-0800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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les 12; Conserv
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TOPOLOGY: li
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TYPE: nucleic acid
STRANDEDNESS: single
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TAKAHA, Takeshi
YANASE, Michiyo
OKADA, Shigetaka
                                                                                                                                                                                                                                                                                                                                         NAKAMURA, Hiroyasu
                                                                                                                                                                                                                                                                                                                       FUJII, Kazutosh
                                                                                                                                                                                                                                                                                                                                                         TAKATA, Hiroki
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100.0%; F1
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US-09-221-017B-225
                                                                 US-09-221-017B-225
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                                Query Match
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PP1182
FILING DATE: 31-DEC-1997
PRIOR APPLICATION DATA:
APPLICATION UMBER: PP1546
FILING DATE: 30-JAN-1998
PRIOR APPLICATION DATA:
APPLICATION UMBER: PP2911
FILING DATE: 09-APR-1998
PRIOR APPLICATION UMBER: PC7/AU98/01023
APPLICATION NUMBER: PC7/AU98/01023
FILING DATE: 10-DEC-1998
FILING DATE: 10-DEC-1998
APPLICATION NUMBER: PC7/AU98/01023
FILING DATE: 10-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM COMPALIALE
OPERATING SYSTEM: Windows
SOFTWARE: FASTSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/221,017B
APPLICATION NUMBER: 21-DEC-1998
                                                                           FEATURE:
NAME/KEY: misc_feature
LOCATION: 1...2572
                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 2572 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                ANTI-SENSE: UNFORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                             HYPOTHETICAL:
                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                        TELEFAX: 000141
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ITLE OF INVENTION: P. GINGIVALIS NUCLEOTIDES AND USES THEREOF
Local Similarity
nes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                             NAME: Monroy, Gladys H
REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                  ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94304-1018
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ilarity 100.0%;
Conservative (
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                                                                                                                                  PORYPHYROMONAS GINGIVALIS
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               57.1%; Score 12; 100.0%; Pred. No.
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                 DB 4; Length 2572;
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0; Indels
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RESULT 73
US-09-221-017B-462/c
; Sequence 462, Application US/09221017B
; Patent No. 6444799
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; LOCATION: (4)..(2592)
US-09-619-353-13
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SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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CURRENT APPLICATION NUMBER: US/09/619,353
CURRENT FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 60/144,766
PRIOR FILING DATE: 1999-07-20
NUMBER OF SEQ ID NOS: 14
SOFTWARF: DATE: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION: APPLICANT: Ross,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Brachydanio rerio (zebrafish)
                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: PP1182
FILING DATE: 31-DEC-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                        ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ROSS, Bruce C.
TITLE OF INVENTION: P. GII
NUMBER OF SEQUENCES: 1120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TLE OF INVENTION: Odorant Receptors
                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
APPLICATION NUMBER: PP1546
FILING DATE: 30-JAN-1998
RIOR APPLICATION DATA:
                                                                                                             FILING DATE: 2
CLASSIFICATION:
                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                        CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                      STREET:
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No. 6410249
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Dittman, Andrew H.
Fan, Jinhong
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09619353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lin, David M.
                                                                                                                                                                                                                                                                                                                         755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                               23-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                              Bruce C.
N: P. GINGIVALIS NUCLEOTIDES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57.1%; Score 12;
100.0%; Pred. No.
                                                                                                                                              US/09/221,017B
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. 95;
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FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR PILING DATE: 1998-02-18

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR PILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 3523

LENGTH: 2832
RESULT 75
US-08-936-165A-66/c
; Sequence 66, Application US/08936165A
; Patent No. 6348582
                                                                                                           밁
                                                                                                                                                                                                                                                  ; TYPE: DNA
, ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-3523
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US-09-252-991A-3523
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6551795
GENERAL INFORMATION:
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                                                                                                                                                                              Matches
                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3523, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELAT
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 2760 base pairs
TYPE: nucleic acid
STRANDENNESS: double
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 10-DEC-1998
ATTORNEY/AGENT INFORMATION:
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Local Similarity 100.0%; P
hes 12; Conservative 0;
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REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
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                                                                                                       1490 CGTTCGAGATGA 1501
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                                                                                                                                          9 CGTTCGAGATGA 20
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NUCLEIC ACID AMD AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                               57.1%; Score 12; 100.0%; Pred. No.
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95;
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INFORMATION FOR SEQ ID NO: 66:
SEQUENCE CHARACTERISTICS:
LENGTH: 2854 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
US-08-936-165A-66
                                                                                      밁
Search completed: December 17, 2003, 10:01:51 Job time : 67 secs
                                                                                                                                                                   Query Match 57.1%; Score 12; DB 4; Length 2854; Best Local Similarity 100.0%; Pred. No. 95; Matches 12; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936,165A
FILING DATE: 24-SEP-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/027,032
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 24-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Gimmi, Edward R
REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: P50549
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
TELEPHONE: 610-270-5090
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APPLICANT: Black,
APPLICANT: Burnhau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Pratt, Julie
APPLICANT: Pratt, Julie
APPLICANT: Reichard, Richard
APPLICANT: Resemberg, Martin
APPLICANT: Ward, Judith
TITLE OF INVENTION: No. 6348582el Prokaryotic Polynucleotides,
TITLE OF INVENTION: Polypeptides and Their Uses
NUMBER OF SEQUENCES: 534
CORRESPONDENCE ADDRESS:
ADDRESSEB: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 709 Swedeland CITY: King of Prussia
                                                                                    1984 CGTTCGÁGÁTGÁ 1973
                                                                                                              9 CGTTCGAGATGA 20
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Hodgson, John
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Lonetto, Michael
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